MULTIRESIDUE METHODS FOR THE DETERMINATION OF CHLORINATED PESTICIDES AND POLYCHLORINATED BIPHENYLS (PCBS) IN WILDLIFE TISSUES BY GAS CHROMATOGRAPHY/MASS SPECTROMETRY

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PREFACE

The Canadian Wildlife Service's National Wildlife Research Centre (NWRC) has been providing analytical determinations of chlorinated pesticides and polychlorinated biphenyl (PCB) congeners in wildlife tissues since 1986, in support to its National Wildlife Toxicology Program. Starting in 1997, all analytical data were determined using a quadrupole mass selective detector (MSD) coupled to the gas chromatograph instead of the electron capture detector (ECD) system that had been used up to that time. This change has allowed the unequivocal identification of a larger number of compounds.

This report provides a detailed description of the GC/MSD methodology along with information on analytical quality control. The first method, "MET-CHEM-OC-04", which describes the analysis of various biological matrices, is followed by "MET-CHEM-OC-03" which is applicable more specifically to adipose tissues and milk from the polar bear. The majority of the methodology used for the polar bear tissue is the same as that used for other tissues, the differences are mainly in the identification of particular compounds which are more commonly determined in the polar bear tissue.

Standard Operating Procedures (SOPs) specific to the organization are cited throughout this document. These SOPs are not readily available in the published literature but can be obtained from the authors, upon request

The names of manufacturers, suppliers and trade names are included only to document the exact assay conditions adopted by the NWRC. Other equivalent products, instruments or reagents from other sources may also give satisfactory results.

PRÉFACE

Méthodes multi-résidus pour le dosage des pesticides organo-chlorés et des biphényles polychlorés (BPC) dans des tissus d'espèces sauvages.

Depuis 1986, le Centre national de la recherche faunique (CNRF) du Service canadien de la faune fournit des résultats d'analyse de pesticides organo-chlorés et de congénères de biphényles polychlorés (BPC), pour appuyer les projets de recherche reliés à son Programme national de surveillance des effets des produits toxiques sur les espèces sauvages. Depuis 1997, les résultats d'analyse sont obtenus utilisant un appareil de spectrométrie de masse de type quadripolaire couplé au chromatographe en phase gazeuse (GC/MSD), plutôt qu'un système de détection à capture d'électron (GC/ECD). Cette modification a permis l'identification unéquivoque d'un plus grand nombre de composés.

Le présent document donne une description détaillée de la méthode améliorée (GC/MSD) ainsi que l'information reliée aux activités de contrôle de qualité. La première méthode « MET-CHEM-OC-04 », décrivant l'analyse de différentes matrices d'origine biologique, est suivie de la méthode « MET-CHEM-OC-03 », applicable plus spécifiquement aux tissus adipeux et au lait d'ours polaire. Les étapes de la méthode pour l'analyse des tissus d'ours polaire sont très semblables à la méthode utilisée pour les autres tissus, les principales différences étant surtout reliées à l'identification des composés que l'on retrouve plus fréquemment dans les tissus d'ours polaires.

Tout au long du document on fait référence à des modes opératoires normalisés (« SOPs ») qui sont spécifiques à notre organisation. Ces procédures ne sont pas disponibles dans la littérature mais peuvent être obtenues en communiquant directement avec les auteurs.

Le nom des manufacturiers, fournisseurs et nom de commerce des produits sont inclus uniquement dans le but de documenter les conditions d'analyse précises utilisées par le CNRF. Des produits, instruments ou réactifs équivalents provenant d'autres sources peuvent aussi donner des résultats satisfaisants.

ABBREVIATIONS

DCM dichloromethane

GC gas chromatograph

GPC gel permeation chromatography

HGQA Herring gull eggs quality assurance material

MDC minimum detectable concentration (3x noise level)

MSD mass selective detector

MS mass spectrometer

OC organochlorine

PCB polychlorinated biphenyl

PFTBA perfluorotributylamine

SIM selected ion monitoring

SRM standard reference material

TCPM tris(4-chlorophenyl) methanol

ACKNOWLEDGEMENTS

The authors gratefully acknowledge the outstanding work of Ghislaine Sans Cartier in the preparation of this report. She has provided vital skills in editing data and text, collating information from the authors and has been instrumental in developing a clear layout of the material. We also wish to thank John Moisey and Mary Simon for the technical review of this document.

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MET-CHEM-OC-03 /

MULTIRESIDUE METHOD FOR THE DETERMINATION OF CHLORINATED PESTICIDES AND POLYCHLORINATED BIPHENYLS (PCBs) IN POLAR BEAR TISSUES

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MULTIRESIDUE METHOD FOR THE DETERMINATION OF CHLORINATED PESTICIDES AND POLYCHLORINATED BIPHENYLS (PCBs) IN WILDLIFE TISSUES

1. Scope and Field of Application

The method that follows is applicable to the analysis of various matrices such as egg, liver, breast muscle and plasma for the determination of trace levels of chlorinated pesticides (OCs) and polychlorinated biphenyls (PCBs). There are specific contaminants which are usually targeted and they are listed in **Tables 6** and 7. Typical detection limits of 0.001 ppm (or less) are obtained for most compounds of interest. For the analysis of tissues and milk from polar bear (*Ursus maritimus*), a description of the methodology used is included in the second part of this report (method "MET-CHEM-OC-03").

2. REFERENCES

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3. PRINCIPLES AND DEFINITIONS

This method is intended for the analysis of lipophilic trace organic chemical contaminants in biological tissues. The first step, to extract the lipids from the tissue, is accomplished by extracting dried tissue homogenates with DCM/hexane or, in the case of plasma, by first denaturing it with formic acid prior to the DCM/hexane extraction. Chemicals of interest in the extracts are then separated from lipids and biogenic compounds by gel permeation chromatography (GPC), except for plasma extracts for which it is unnecessary. Any residual lipids are removed by Florisil column chromatography. Finally, the OCs and PCBs levels are determined via high resolution gas chromatography coupled to a mass spectrometry detection system. Identification of target analytes is accomplished by comparing GC retention times and specific mass fragments known to be present in the spectra of authentic compounds. Quantitation is accomplished by comparing the intensity of targeted mass fragments in specimen extracts to the same compounds in standard mixtures, injected separately on the GC/MSD system.

4. REAGENTS, SOLUTIONS AND STANDARDS

SAFETY PRECAUTIONS

- ⇒ The toxicity or carcinogenicity of each reagent and standard used in this method has not been precisely defined. Each chemical must be treated as a potential health hazard and exposure should be reduced to the lowest possible level.
- ⇒ Operations with toluene, DCM, hexane, methanol and acetone should be performed in a fume hood and dermal contact with solvents should be avoided.
- ⇒ Standards should always be opened and used in a fume hood. Handling of these compounds must be done only by qualified technical staff.
- ⇒ General safety rules and waste disposal procedures that apply to the Trace Organic Chemistry Laboratory must be followed (ref. Safety Manual).
- ⇒ Material Safety Data Sheets (MSDSs) for the products used in the assay must be read.

4.1. Reagents

- 4.1.1. Acetone, Omnisolv®, EM Science AX0116-1
- 4.1.2. Hexane, Omnisolv®, EM Science HX0296-1
- 4.1.3. Dichloromethane, Omnisolv®, EM Science DX0831-1
- 4.1.4. Methanol, Omnisolv®, EM Science MX0488-1
- 4.1.5. Toluene, Omnisolv®, EM Science TX0737-1
- 4.1.6. 2,2,4-Trimethylpentane (iso-octane), Omnisolv®, EM Science TX1389-1
- 4.1.7. Formic acid, AnalaR®, EM Science B10115
- 4.1.8. o-Phosphoric acid, 85%, HPLC grade, Fisher Scientific A260-500
- 4.1.9. Sulfuric acid, Fisher Scientific A510-500
- 4.1.10. Sodium sulfate, anhydrous granular (Na₂SO₄), EM Science ACS850-46

Wash 600 g of Na_2SO_4 in a glass column 3 cm ID x 50 cm long with 600 mL DCM/hexane (1:1), air dry in an open dish under the fume hood, heat 3 h at 400°C in a muffle furnace, cool and transfer in a tightly capped glass bottle. *Note:* If, after

heating, the sodium sulfate develops a grayish cast (due to the presence of carbon in the crystal matrix), discard that batch.

- 4.1.11. Ethyl alcohol, distilled in glass, EM Science EX0278-6
- 4.1.12. Vanillin, crystalline, Fisher Scientific V10-100
- **4.1.13.** Olive oil, Acros Organic 41653-5000
- 4.1.14. De-ionized water from the Milli-RO / Milli-Q system (Millipore)
- 4.1.15. Helium, compressed bottled gas, Praxair, HE UHP SG 103168K
- 4.1.16. Nitrogen, compressed bottled gas, Praxair, N₂ PRE PURE SG 105411K

4.2. Adsorbents for Sample Cleanup

- 4.2.1. Envirobeads™ S-X3, Select (200-400 mesh), ATS Scientific 091-203
- 4.2.2. Florisil®, pesticide grade, 60-100 mesh, EM Science B28722-44

Transfer 750 to 800 g of Florisil into a large porcelain dish and heat to 650°C for 6 h in a muffle furnace. Reduce furnace to 130°C and heat overnight. Transfer a known amount of Florisil to a 4 L amber glass bottle. Cool with lid closed. Add 1.2% deionized water to the Florisil. Close bottle with Teflon lined lid and shake contents to break up lumps. Transfer to a roller and agitate for 24 h to equilibrate before use. Shelf life: 2 months. If a longer period has elapsed since preparation, dry Florisil overnight at 130°C, cool in closed container, add de-ionized water and equilibrate as described above.

4.3. Solutions

- **4.3.1.** DCM/hexane, 1:1 (v/v)
- **4.3.2.** Vanillin reagent, 6 g/L Dissolve 6.0 g of vanillin in water in a 1 litre volumetric flask and dilute to volume. Transfer in amber bottle and store at room temperature. **Shelf life:** stable for two months.
- **4.3.3.** Phospho-vanillin reagent Add 350 mL of vanillin reagent (4.3.2) and 50 mL of water to a 2 litre beaker. Add 600 mL of concentrated ophosphoric acid, with constant stirring. Transfer in amber bottle and store at room temperature. **Shelf life:** stable for two months.

4.4. Stock Standards

Note: Details are given in **Table 1** and list of PCB congeners (based on Ballschmiter's nomenclature) is given in **Appendix A**.

- **4.4.1.** Isotopically-labeled 13 C₁₂-PCBs prepared in n-Nonane Cambridge Isotope Laboratories (CIL), certified solution: $40 \pm 4 \mu g/mL$
- 4.4.2. A mixture of isotopically-labeled 13 C₆-chlorobenzene prepared in isototane Cambridge Isotope Laboratories (CIL) certified solution: $100 \pm 10 \,\mu\text{g/mL}$
- **4.4.3.** Native (unlabeled) organochlorines solid compounds from various suppliers. Stock solutions prepared in iso-octane as described in 4.5.1
- **4.4.4.** Aroclor 1242, Aroclor 1254 and Aroclor 1260 US EPA and Monsanto. Stock solution (with the three Aroclors) prepared in iso-octane as described in 4.5.2

4.5. Intermediate Standard Solutions

Note: Refer to SOP-CHEM-PROC-05 for details concerning the preparation and storage of standard solutions.

- 4.5.1. Organochlorines Accurately weigh about 10 mg (between 10 to 30 mg, depending on the compound) of each of the twenty-one organochlorines quantitative standards listed in Table 1. Dissolve with iso-octane in the same 100 mL volumetric flask and dilute to volume (final concentration for each compound varies from 100 to 300 μg/mL). Store in 125 mL Hypo-vial and cap with Mininert valve at 4°C.
 - *Note:* Hexachlorobenzene and nonachlor need to be dissolved in a small amount of toluene before dilution with iso-octane.
- 4.5.2. Aroclors Accurately weigh about 1 g of each of the three Aroclor standards listed in **Table 1**. Dissolve with iso-octane in the same 100 mL volumetric flask and dilute to volume. Store in 125 mL Hypo-vial and cap with Mininert valve at 4°C.

4.6. Working Standard Solutions

4.6.1. Internal standard spiking solution - Add 500 μL of each of the 6 ¹³C₁₂-PCBs and 250 μL of the chlorobenzene cocktail (ref. **Table 1**) in a 10 mL volumetric flask and dilute to volume with toluene.

- **4.6.2.** Normalization standard Dilute 3 mL of the ¹³C₁₂-PCB-138 solution in 27 mL toluene. Note: Normalizing on area responses of the ¹³C₁₂-PCB-138 in samples and standards corrects for changes in the MSD sensitivity between injections and changes in the final sample volumes.
- **4.6.3.** Quantitation standards -
 - **4.6.3.1.** Organochlorines Pipette 3 mL of the solution containing 21 OCs (Section 4.5.1) in a 1 L volumetric flask and dilute to volume with iso-octane. Mix well and store at room temperature away from light and heat.
 - 4.6.3.2. Aroclors (PCBs) Pipette 1 mL of the solution of Aroclors (Section 4.5.2) to a 100 mL volumetric flask and dilute to volume with iso-octane. Mix well and store at room temperature away from light and heat. This solution contains 100 μg/mL of each Aroclors (i.e., 0.3 μg/μL total PCBs). Working standard is prepared by diluting 3 mL of this solution to 100 mL with iso-octane. The final solution concentration will be 9000 pg/μL total PCBs.
- 4.6.4. Initial calibration standards Calibration standards (CS1 through CS5) for OCs are prepared by diluting the solution prepared in Section 4.5.1 in iso-octane, to produce the concentrations in the working range shown in Table 2. For the PCBs, calibration standards (CS1 through CS5) are prepared by diluting the solutions in Section 4.5.2 to obtain concentrations shown in Table 3, ranging from 900 pg/μL to 30 000 pg/μL total PCBs. Note: PCB congeners concentrations shown are based on actual quantitation of each congener in the Aroclor solutions used [2.2].

4.7. Mass Spectrometer Calibration Standard

Perfluorotributylamine (PFTBA), cat. no. 110234-0, PCR Inc., Gainesville, FL.

4.8. Olive Oil Standard

Note: Used for plasma lipid determination.

- **4.8.1.** Add ca 150 mg of olive oil to a 10 mL volumetric flask and dilute to volume with ethyl alcohol. Mix well and store at 4°C. **Shelf life:** one month.
- 4.8.2. For each set of assays, prepare 4 calibration standards to have

concentration ranging from 30 to 150 mg/10 mL by diluting the olive oil stock solution in ethyl alcohol.

4.9. QA Reference Material

Herring gull (*Larus argentatus*) egg homogenate prepared in-house from eggs collected in 1989 from Lake Ontario and diluted with domestic chicken eggs, to adjust contaminant concentrations. *Note:* Details on the preparation of this quality assurance material is given in Wakeford 1997 [2.1].

4.10. Method Blank

Clean reference matrix carried out through the entire analytical procedure. **Note:** chicken egg is used when contaminant free material with the same matrix as the sample is not available.

TABLE 1 - Supplier, catalogue number and concentrations of PCBs and organochlorines standards

Inter	nal Standards			Solution o	concentrations
		Supplier	Cat. no.	Std. (µg/mL)	Working Std. (ng/µL)
3 C 12-1	PCB				
<u> </u>	¹³ C ₁₂ -PCB-28	CIL	EC-1413	40	2
	¹³ C ₁₂ -PCB-52	CIL	EC-1424	40	2
	¹³ C ₁₂ -PCB-118	CIL	EC-1435	40	2
	¹³ C ₁₂ -PCB-153	CIL	EC-1406	40	2
	¹³ C ₁₂ -PCB-180	CIL	EC-1407	40	2
	¹³ C ₁₂ -PCB-194	CIL	EC-1418	40	2
				40	
3 C 6-C	Chlorobenzene cocktail containing:	CIL	EM-1725-A		
	13C ₆ -1,2,4,5-tetrachlorobenzene			100	2.5
	13C ₆ -1,2,4,5-pentachlorobenzene			100	2.5
,	¹³ C ₆ -1,2,4,5-hexachlorobenzene			100	2.5
Norm	nalization Standard				
	¹³ C ₁₂ -PCB-138	CIL	EC-1436	40	4
	<u></u>			***************************************	
Quar	ntitation Standards				
Organ	ochlorines			Std. (µg/mL)	Working Std. (pg/µL
	1,2,4,5-tetrachlorobenzene	Ultra Scientific	RCP-29	100	300
	1,2,3,4-tetrachlorobenzene	Ultra Scientific	RCP-27	100	300
	pentachlorobenzene	Ultra Scientific	RCP-30	100	300
	hexachlorobenzene	US EPA	3920	100	300
	alpha-hexachlorocyclohexane	US EPA	0620	100	300
	beta-hexachlorocyclohexane	US EPA	0640	100	300
	gamma-hexachlorocyclohexane	US EPA	0680	100	300
	octachlorostyrene	private source	n/a	100	300
	oxychlordane	US EPA	5200	300	900
	trans-chlordane	US EPA	1240	200	600
	cis-chlordane	US EPA	1220	200	600
	trans-nonachlor	US EPA	5080	200	600
	cis-nonachlor	private source	n/a	100	300
	p,p'-DDE	US EPA	1860	300	900
	p,p'-DDD	US EPA US EPA	1780 1920	300	900
	p,p'-DDT	private source	1920 n/a	100	300
	photo-mirex mirex	US EPA	n/a 4720	200	600
	heptachlor epoxide	US EPA	3880	100	300
	dieldrin	US EPA	2380	200	600
	tris(4-chlorophenyl)methanol	Lancaster	n/a	100	300
PCBs	the(4-onto option)) mentano	- COLICOSTEI	1 IV CI	100	300
, ,,,,	Aroclor-1242	US EPA	5703	10000	3000
	Aroclor-1254	Monsanto	n/a	10000	3000
	Aroclor-1260	Monsanto	n/a	10000	3000

Chemistry Unit	
Laboratory Services Section	n. NWRC

CWS Technical Report No. 335 Method No. MET-CHEM-OC-04

TABLE 2 - Composition of OCs initial calibration standards (CS) in pg/ μL

Organochiorines	CS1	CS2	CS3 °	CS4	CS5
1,2,4,5-tetrachlorobenzene	30	100	300	500	1000
1,2,3,4-tetrachlorobenzene	30	100	300	500	1000
pentachlorobenzene	30	100	300	500	1000
hexachlorobenzene	30	100	300	500	1000
alpha-hexachlorocyclohexane	30	100	300	500	1000
beta-hexachlorocyclohexane	30	100	300	500	1000
gamma-hexachlorocyclohexane	30	100	300	500	1000
octachlorostyrene	30	100	300	500	1000
oxychlordane	90	300	900	1500	3000
trans-chiordane	60	200	600	1000	2000
cis-chlordane	60	200	600	1000	2000
trans-nonachlor	60	200	600	1000	2000
cis-nonachlor	30	100	300	500	1000
p,p'-DDE	90	300	900	1500	3000
p,p'-DDD	90	300	900	1500	3000
ρ,p'-DDT	90	300	900	1500	3000
photo-mirex	30	100	300	500	1000
mirex	60	200	600	1000	2000
heptachlor epoxide	30	100	300	500	1000
dieldrin	60	200	600	1000	2000
tris(4-chlorophenyl)methanol	30	100	300	500	1000

a used daily to verify calibration and abundance ratios

TABLE 3 - Composition of PCB congeners initial calibration standards (CS) in $pg/\mu L$

PCBs		CS1	CS2	CS3 °	CS4	CS5
	PC8-18	24	81	244	406	812
	PCB-17	16	53	159	265	530
-	PCB-16/32	16	52	156	259	519
	PCB-31	19	65	195	324	649
	PCB-28	21	70	209	348	695
	PC8-33/20	19	63	188	314	627
	PCB-22	9	29	87	145	290
	PCB-52	28	92	275	458	917
	PCB-49	12	40	120	200	400
	PCB-47/48	7	25	74	124	247
	PCB-44	17	58	173	289	577
		11	36	107	178	357
	PC8-42	13	44	132	221	442
	PC8-64				143	
	PCB-74	9	29	86	363	285 725
	PCB-70/76	22	73	218		
	PCB-95	16	54	163	272	544
	PCB-66	24	82	245	408	816
	PC8-56/60	11	35	105	176	352
	PCB-92	6	19	57	95	189
	PCB-101/90	36	121	362	604	1208
	PCB-99	11	38	113	188	377
	PCB-97	9	31	94	157	314 ⁻
	PCB-87	16	54	163	272	544
	PC8-85	5	17	51	85	170
	PCB-110	32	108	323	538	1076
	PCB-151	16	55	164	274	548
	PCB-149	37	124	371	619	1238
		1				
	PCB-118 ^b	23	78	235	391	782
	PC8-146	5	18	55	91	182
	PCB-153	40	133	400	666	1333
	PCB-105	12	41	122	204	408
	PCB-179	7	25	75	125	249
	PCB-141	11	36	107	178	356
	PCB-130	1	- 5	14	24	48
	PC8-176	2	8	25	41	82
	PCB-137	2	6	· 19	32	65
	PCB-158	5	16	48	80	160
	PCB-138	44	146	438	730	1460
	PCB-178	4	15	44	73	147
	PCB-187	18	59	177	296	591
	PCB-183	9	29	86	144	287
	PCB-128	6	19	56	93	187
	PCB-174	17	55	166	277	555
	PC8-177	9	30	90	150	299
	PC8-202	2	7	22	36	72
				38	64	128
	PCB-171	4	13		l	
	PCB-156	6	18	55	92	184
	PCB-200	2	6	19	31	62
	PC8-157	11	4	13	22	45
	PCB-172	3	8	25	42	83
	PCB-180	33	109	328	547	1093
	PCB-170/190	17	55	166	276	552
	PCB-201	8	26	78	130	260
	PCB-196/203	9	29	87	145	290
	PC8-208	2	6	17	29	57
	PCB-195	3	10	31	52	104
	PCB-207	2	5	16	27	54
	PC8-194	8	25	75	126	251
	PCB-206	2	7	22	36	73

a used daily to verify calibration and abundance ratios

^b calibration curve shown in Figure 3

5. AUXILIARY EQUIPMENT

5.1. Glassware and Labware

- **5.1.1.** Aluminum foil
- 5.1.2. Amber glass bottle 1 L and 4 L with Teflon lined screwed cap
- **5.1.3.** Autosampler vials 2 mL, clear with red/orange crimp caps (HP 5181-3400)
- 5.1.4. C18 cartridge (Superclean ENVI-18, 6 mL tubes) Supelco 505706
- **5.1.5.** Centrifuge tubes, graduated, glass 15 mL with \$13 ground glass stopper
- 5.1.6. Centrifuge tubes, glass 50 mL with screw-top
- 5.1.7. Column, glass, 1.0 cm ID x 24 cm long with Teflon[™] stopcock and with \$19/22 outer joint at top of column (used for Florisil cleanup)
- 5.1.8. Column, glass 2.1 cm ID x 35 cm long with Teflon™ stopcock and with \$24/40 outer joint at top of column (used for sample extraction)
- 5.1.9. Column glass 3 cm ID x 50 cm long with Teflon™ stopcock and reservoir (for preparing the Na₂SO₄)
- **5.1.10.** Cuvettes, acrylic, disposable (Sarstedt 67.738)
- 5.1.11. Flasks, flat bottom 125, 250 and 500 mL all with \$24/40 outer joint
- 5.1.12. Flasks, pear shaped 5 and 10 mL
- 5.1.13. Funnel, glass 25 mm and 10 cm
- **5.1.14.** Glass wool (Canadawide Scientific 54100-11), pre-washed with DCM/hexane (1:1) and air dried
- 5.1.15. GPC glass column 3 cm ID x 60 cm long Envirosep-ABC column assay (ABC Laboratories Inc., Columbia, MO, USA)
- 5.1.16. GPC tubes, 20 mL (ABC Laboratories Inc., Columbia, MO, USA)
- 5.1.17. Graduated cylinders, glass 10, 50, 100 and 500 mL, and 1 L
- 5.1.18. Hand crimper, 8 mm for crimping aluminum seals to autosampler vials (Chromatographic Specialties)
- 5.1.19. Mortars and pestles, glass
- 5.1.20. Pasteur pipets
- 5.1.21. Pipet, Eppendorf 5-100 μ L (with tips)

- **5.1.22.** Pipets, glass, disposable 0.5, 1, 2, 5 and 10 mL
- 5.1.23. Pipet, Rainin, with tips
- 5.1.24. Porcelain dish 21.5 cm diameter
- **5.1.25.** Reacti-vials 1 and 5 mL and Hypo-vials 125 mL with MininertTM valve (Chromatographic Specialties Inc.)
- **5.1.26.** Reservoir, glass 125 mL, with \$19/22 inner joint for glass column 5.1.7
- 5.1.27. Reservoir, glass 250 mL, with \$24/40 inner joint for glass column 5.1.8
- **5.1.28.** Spatulas
- **5.1.29.** Syringe, glass 10 mL (B-D D3037)
- 5.1.30. Syringes, HamiltonTM 10, 50, 100 and 250 µL
- 5.1.31. Volumetric flasks, glass 10, 50 and 100 mL, and 1 L
- 5.1.32. Weighing aluminum dishes disposable

5.2. Equipment

- **5.2.1.** Analytical balance (Sartorius BP210D) and top-loading balance (Mettler PR700)
- 5.2.2. Vortex mixer
- **5.2.3.** Rotary evaporator with water bath at ca 30°C (Büchi Rotavapor-R, Brinkman Instruments)
- 5.2.4. Refrigerated circulating bath at ca -4°C (Lauda RM20)
- 5.2.5. Drying oven (Fisher Scientific, Model 516 G)
- 5.2.6. Muffle furnace (Blue M Electric Company, Blue Island, Il, USA)
- 5.2.7. Centrifuge (Sorvall, SPX) with Sorvall Powerstat (Type 108-1164)
- **5.2.8.** Visiprep solid phase extraction vacuum manifold (Supelco 57030) with Visidry drying attachment (Supelco 57100)
- 5.2.9. Water bath at 100°C (Blue M) for plasma lipids
- 5.2.10. Water bath at 37°C (Precision Instruments) for plasma lipids
- **5.2.11.** Roller culture apparatus (Wheaton Instruments)

5.3. Instrumentation

- 5.3.1. GPC Autoprep[™]1000, with automatic sample loading and sample collection unit (from O.I. Analytical), with 23 sample ports (10 mL volume).
- 5.3.2. GC/MSD, Hewlett-Packard gas chromatograph (GC) 5890 Series II equipped with an autosampler (7673A), a Galileo Channeltron electron multiplier (5778) and linked to a Hewlett-Packard 5970 (or 5971A) mass selective detector (MSD) with MS ChemStation (HP G1034C, Rev. C.02.00); GC column: 30 m DB-5 (J&W) fused silica column, 0.25 mm ID, 0.25 μm film thickness (Chromatographic Specialties J1225032).
- **5.3.3.** Spectrophotometer (Hewlett-Packard Diode Array, Model 8452A) for plasma samples.

6. SPECIMEN OR SAMPLE HANDLING REQUIREMENTS

Samples provided to the Trace Organic Chemistry Laboratory are prepared as described in the Tissue Preparation Unit's standard operating procedure SOP-TP-PROC-07. These tissues were usually collected and preserved as recommended in the document "Protocol for Field Collection and Storage of Wild Birds for Biomarker Studies" (S. Trudeau, Biomarker Laboratory, NWRC, 1992).

7. PROCEDURE

Flow chart that summarize procedures for sample extraction, cleanup and analysis is given in Figure 1.

7.1. Columns Preparation

7.1.1. GPC Column

Place 60 g Envirobeads TM S-X3 in a 500 mL beaker. Cover the beads with DCM/hexane (1:1) and allow to swell overnight (a minimum of 12 h). Pack a GPC column (Section 5.1.15) with the pre-swelled beads.

Note: This material generally makes 43 to 45 cm in column length.

7.1.2. Florisil column

Prepare fresh daily as described in Sections 7.5.1.-7.5.3.

7.2. Extraction - Eggs and tissue samples

Note: For plasma samples see Section 7.3.

- 7.2.1. Grind between 1.5 g to 3.0 g of the homogenized sample with ca 25 g of the treated anhydrous Na₂SO₄ (Section 4.1.10) in a glass mortar and pestle until a free-flowing mixture is obtained. *Note:* A sample size of 3 g is preferred. Include at least one standard reference material and a method blank sample with each batch of samples (a typical batch contains 10 samples).
- 7.2.2. Plug a 2.1 cm ID x 35 cm long glass column (Section 5.1.8) with some treated glass wool (Section 5.1.14), add about 1 cm Na₂SO₄. Pour ground sample mixture into the glass column and tap the column gently to settle the mixture. Add ca 0.5 cm Na₂SO₄. *Note:* Usually 5 columns are prepared at one time.
- 7.2.3. Place a 500 mL flat bottom evaporating flask under the column and open the stopcock. Rinse the mortar and pestle with DCM/hexane (1:1), and transfer the rinse onto the top of the column, repeat rinsing mortar and pestle three times. Close the stopcock.
- 7.2.4. Add enough DCM/hexane to cover the mixture and allow to soak for 30 min.
- 7.2.5. Attach the reservoir (Section 5.1.27) to the column and add 200 mL DCM/hexane (1:1). Elute at 5-10 mL/min (collecting the eluate in the 500 mL evaporating flask).
- 7.2.6. Evaporate the eluate to less than 5 mL on a rotary evaporator with water bath at ca 30°C, then quantitatively transfer into a 15 mL graduated centrifuge tube. Rinse the flask three times with ca 1 mL of DCM/hexane (1:1), adding the rinses to the centrifuge tube.
- 7.2.7. Add DCM/hexane (1:1) to have a sample concentration of 0.2 g/mL. For example with 3 g of tissue, adjust volume to 15 mL.
- 7.2.8. Transfer an aliquot equivalent to 0.5 g tissue into a pre-weighed aluminum dish, to be used for lipid determination (ref. Section 7.14).

7.2.9. Transfer an aliquot equivalent of 1 g of tissue into a GPC tube (Section 5.1.16). Spike the extract with 50 µL of the ¹³C-labeled chlorobenzene/ PCB internal standard spiking solution (Section 4.6.1), and dilute to 10 mL with DCM/hexane (1:1). The sample is now ready for GPC cleanup.

7.3. Extraction - Plasma samples

- 7.3.1. Accurately weigh ca 3 g aliquot of the thawed plasma into a 50 mL screw-top centrifuge tube. Spike with 50 μL of the ¹³C-labeled chlorobenzene /PCB internal standard spiking solution (Section 4.6.1). Mix the spiked plasma gently with a Vortex mixer, and let it stand for 30 min to equilibrate.
- 7.3.2. Add an equal volume of formic acid (e.g., 3 mL for 3 g plasma) to the spiked plasma in order to denature proteins; mix gently with Vortex mixer, and let it stand for 15 min.
- 7.3.3. Activate a C18 cartridge with two 6 mL portions of methanol followed by two 6 mL portions of de-ionized water, using the Visiprep solid phase vacuum manifold (Section 5.2.8).
- 7.3.4. Load the sample mixture (from 7.3.2) onto the activated C18 cartridge with suction at a flow rate of 6-7 mL/min. The polar interferences and lipids are not retained by the cartridge.
- 7.3.5. Dry the C18 cartridge thoroughly with a stream of nitrogen gas using a Visidry Drying attachment for ca 35 min. *Note:* Incomplete drying of the cartridge would result in sample loss.
- 7.3.6. Elute the analytes from the dried C18 cartridge with 3 x 2 mL of DCM/hexane (1:1). The sample is now ready for florisil cleanup (Section 7.6).

7.4. Sample Cleanup by GPC

Note: Calibrate GPC system when the column is changed, when channeling occurs, when column drying has occurred or when recoveries are not acceptable. Refer to the GPC operating manual [2.8] for specific instructions.

7.4.1. Load the GPC tubes (from 7.2.9) into the instrument.

- 7.4.2. Set GPC flow-rate at 5 mL/min of DCM/hexane (1:1), the "dump" time to 26 min, the "collect" time to 36 min and the wash time to 4 min. Initiate the operation of the GPC. *Note:* It is possible to load and run as many as 23 samples simultaneously (typically 12 are used). The sequence can be run overnight.
- **7.4.3.** Evaporate the eluate from 7.4.2 to ca 3 mL on a rotary evaporator. The sample is now ready for Florisil column cleanup.

7.5. Sample Cleanup by Florisil Column

Florisil column cleanup is designed to isolate compounds of interest from any residual lipids.

- **7.5.1.** Plug a 1 cm ID glass column (Section 5.1.7) with some treated glass wool (Section 5.1.14) and attach reservoir (Section 5.1.26).
- 7.5.2. With the stopcock in the close position, add 40 mL hexane to the reservoir and then 8 g of the de-activated Florisil (Section 4.2.2).
- 7.5.3. Open stopcock, and gently tap side of column to settle adsorbent and add ca 1 cm of Na₂SO₄. Close stopcock when solvent level is still slightly above the Na₂SO₄ layer. Discard rinse solvent.
- 7.5.4. Place 150 mL flat bottom evaporating flask at the end of column.
- 7.5.5. Load extract from 7.4.3 (or 7.3.6) on top of the Florisil column using Pasteur pipet. Rinse the flask 3-4 times with small portions of DCM/hexane (1:1). Add rinses to column, allowing solvent to drain to packing level in between rinses.
- 7.5.6. As the last rinse approaches top of Na₂SO₄ layer, add 95 mL of DCM/hexane (1:1) allow elution to proceed at ca 5 mL/min.
- 7.5.7. Concentrate eluate to less than 3 mL with rotary evaporator and quantitatively transfer into a 10 mL pear shaped flask. Rinse flask three times with 1 mL portions of iso-octane adding rinses to the flask.
- 7.5.8. Further concentrate the eluate to less than 400 µL with rotary evaporator. Quantitatively transfer the sample to an autosampler vial, previously marked at the 570 µL level. Rinse flask with a few drops of iso-octane, adding the rinses to the autosampler vial.

- 7.5.9. Add exactly 20 μ L 13 C₁₂-PCB-138 normalization standard (Section 4.6.2) to the autosampler vial using an Eppendorf pipet. Adjust the final volume to the 570 μ L mark with iso-octane. *Note:* Exact volume is not critical, since normalization standard is used to quantify residue levels
- 7.5.10. Cap vial and vortex until thoroughly mixed. The sample is ready to inject into the GC/MSD (see Section 7.12). *Note:* Each sample is injected twice. The first injection is designed to determine OC levels (see chromatographic windows and selected ions in **Table 4**). The second injection is designed to determine PCB levels (see chromatographic windows and selected ions in **Table 5**).

7.6. GC Operating Conditions

7.6.1. Column

 30 m long DB-5 fused-silica column, 0.25 mm ID, 0.25 μm film thickness

7.6.2. Injection information

•	Injection port temperature	250°C
•	Source	Auto
٠	Location	Front
•	Sample washes	0
•	Sample pumps	5
•	Sample volume	$2 \mu L$
•	Viscosity delay	1 s
•	Solvent washes	5A, 5B
•	Splitless injection	
٠	On column	No
٠	Purge A on	1.5 min

7.6.3. Oven temperature programme

- For OCs: 100°C, hold 3 min; 20°C/min to 180°C; 5°C/min to 300°C
- For PCBs: 100°C, hold 3 min; 20°C/min to 180°C; 2.5°C/min to 300°C

7.6.4. Carrier gas (He)

• Head pressure 5-7.5 psi

7.7. MSD Operating Conditions

7.7.1. MSD conditions

• Electron impact (EI) ionization 70 eV (fixed)

Acquisition mode

selected ion monitoring mode (SIM)

• Dwell time (on each ion)

60 ms

• Source temperature

200°C

· Transfer line temperature

280°C

7.8. Data Analysis Parameters

Calibration settings

Reference window: 10.00%

Non-reference window

5.00%

7.9. Compound Information

Typical retention time, target ion and qualifying ion are given in **Table 6 or 7** for analysis of OCs or PCBs respectively. The list is based on those compounds actually found in wildlife tissues.

TABLE 4 - Chromatographic windows and characteristic ions for OCs analysis

SIM Group	Start time	Dwell time	lons
	(min)	(ms)	
Group 1	7.00	60	214, 216, 222, 224 °
Group 2	9.00	60	250, 252, 258 °, 260
Group 3	11.00	60	183, 219, 284, 286, 292 °, 294
Group 4	13.00	60	238, 268 °, 270, 304 °, 306, 345
Group 5	16.00	60	115, 237, 255, 263, 308, 339, 353, 380, 387
Group 6	17.60	60	237, 339, 373, 375
Group 7	18.40	60	246, 318, 373, 375, 407, 409
Group 8	19.00	60	237, 246, 318, 380
Group 9	20.10	60	165, 235, 338 °, 340, 372 °, 374, 407, 409
Group 10	21.70	60	165, 235, 237, 238, 372, 374
Group 11	23,50	60	237, 272, 406 °, 408
Group 12	26.00	60	139, 251, 311, 346, 442 °, 444

 $^{^{}a}$ ions selected to monitor the 13 C-labeled internal standards. Ion 372 is used to monitor the normalization standard (13 C₁₂-PCB-138)

TABLE 5 - Chromatographic windows and characteristic ions for PCBs analysis

SIM Group	Start time (min)	Dwell time (ms)	lons
Group 1	15.00	60	256, 258
Group 2	18.00	60	256, 258, 268 a, 290, 292, 304 a
Group 3	23.00	60	290, 292, 326, 328
Group 4	27.00	60	326, 328, 338 °, 360, 362, 372 °
Group 5	31.00	60	360, 362, 394, 396, 372 *
Group 6	35.00	60	326, 328, 360, 362, 394, 396, 406 °
Group 7	40.00	60	428, 430, 442°, 464, 498

^a ions selected to monitor the ¹³C-labeled internal standards

TABLE 6 - Retention time and target ions for OC compounds, internal standard and normalization standard

Compound ^a	RT ^b (min)	RT window	Tgt ° (m/z)	QI d	Abundance ratio ^e
Ocs (native)					
1,2,4,5-tetrachlorobenzene	7.73	7.23 - 8.23	216	214	77
1,2,3,4-tetrachlorobenzene	8.23	7,73 - 8.73	216	214	77
pentachlorobenzene	9.77	9.27 - 10.27	250	252	62
hexachlorobenzene	12.16	11.66 - 12.66	284	286	81
alpha-hexachlorocyclohexane	11.91	11.41 - 12.41	219	183	
beta-hexachlorocyclohexane	12.57	12.07 - 13.07	219	183	
gamma-hexachiorocyclohexane	12.78	12.28 - 13.28	219	183	
octachlorostyrene	17.10	16.60 - 17.6	308	380	
heptachlor epoxide	17.29	16.79 - 17.79	353	237	
oxychlordane	17.34	16.84 - 17.84	387	115	
trans-chlordane	18.08	17.58 - 18.58	375	373	88
cis-chlordane	18.59	18,09 - 19,09	375	373	88
trans-nonachlor	18.77	18.27 - 19.27	409	407	88
p,p'-DDE	19.23	18.73 - 19.73	246	318	
dieldrin	19.41	18.91 - 19.91	380	237	
p,p'-DDD	20.82	20,32 - 21,32	235	165	
cis-nonachlor	20.92	20,42 - 21,42	409	407	88
p,p'-DDT	22.01	21,51 - 22.51	235	165	
photomirex	23.65	23,15 - 24,15	238	272	
mirex	25.69	25.19 - 26.19	272	237	
tris(4-chlorophenyl)methanol	28.93	28.43 - 29.43	251	139	
¹³ C-internal standard					
¹³ C ₆ -1,2,4,5-tetrachlorobenzene	7.73	7.23 - 8.23	224	222	77
13C ₆ -1,2,4,5-pentachlorobenzene	9.77	9.27 - 10.27	258	260	65
13C6-1,2,4,5-hexachlorobenzene	12.16	11.66 - 12.66	292	294	82
¹³ C ₁₂ -PCB-28	14.36	13.86 - 14.86	268	270	97
¹³ C ₁₂ -PCB-52	15.40	14.90 - 15.9	304	306	77
¹³ C ₁₂ -PCB-118	20.41	19.91 - 20.91	338	340	65
¹³ C ₁₂ -PCB-153	21.19	20.69 - 21.69	372	374	81
¹³ C ₁₂ -PCB-180	24.52	24.00 - 25.02	406	408	97
¹³ C ₁₂ -PCB-194	27.79	27.29 - 28.29	442	444	88
Normalization standard					
¹³ C ₁₂ -PCB-138	31.90	31.40 - 32.40	372		

^aAbbreviations:

DDE - dichloro diphenyl dichloro ethylene

DDD - dichloro diphenyl dichloro ethane

DDT - dichloro diphenyl trichloro ethane

PCB - polychlorinated biphenyl

^b Retention time. Oven program: 100°C, hold 3 min; 20°C/min to 180°C; 5°C/min to 300°C

^{&#}x27;Target ion - Mass (m/z)

^dQualifying ion

^{&#}x27;Abundance ratio - applicable only to ions within molecular ion clusters

TABLE 7 - Retention time and target ions for PCB compounds

Compound ^a	RT ^b (min)	RT window	Tgt ° (m/z)	QI d	Abundance ratio ^e
PCBs (native)					
PCB-18	16.81	16,48 - 17,48	256	258	97
PCB-17	16.90	16,50 - 17.50	256	258	97
PCB-16/32	17.64	17.30 18.30	256	258	97
PCB-31	18.82	18.58 - 19.58	256	258	97
PCB-28	18.89	18.52 - 19.52	256	258	97
PCB-33/20	19.35	19.04 - 20.04	256	258	97
PCB-22	19.72	19.42 - 20.42	256	258	97
PCB-52	20.58	20,52 - 21.52	292	290	77
PCB-49	20.81	20,52 - 21.52	292	290	77
PCB-47/48	21.00	20.71 - 21.71	292	290	77
PCB-44	21.65	21.37 - 22.37	292	290	77
PCB-42	21.84	21.56 - 22.56	292	290	77
PCB-64	22.32	22.00 23.00	292	290	77
PCB-74	23.60	23,34 - 24.34	292	290	77
PCB-70/76	23.80	23.54 - 24.54	292	290	77
PCB-95	24.06	23.89 - 24.80	326	328	65
PC8-66	24.01	23.75 - 24.75	292	290	77
PCB-56/60	24.98	24.72 - 25.72	292	290	77
PCB-92	25.06	24,80 - 25.80	326	328	65
PCB-101/90	25.40	25.14 - 26.14	326	328	65
PCB-99	25.71	25,46 - 26,46	326	328	65
PCB-97	26.67	26.42 - 27.42	326	328	65
PCB-87	26.98	26.73 - 27.73	326	328	65
PC8-85	27.24	27.00 - 28.00	326	328	65
PCB-110	27.58	27,34 - 28.34	326	328	65
PCB-151	28.36	28.12 - 29.12	360	362	81
PCB-149	29.10	28,87 - 29.87	360	362	81
PCB-118	29.21	28.97 - 29.97	326	328	65
PCB-146	30.27	30.05 - 31.05	360	362	81
PCB-153	30.66	30.43 - 31.43	360	362	81
PCB-105	30.87	30.64 - 31.64	326	328	65
PCB-179	31.58	31.35 - 32.35	394	396	97
PCB-141	31.46	31,23 - 32.23	360	362	81
PCB-130	31.92	31.69 - 32.69	360	362	81
PCB-176	32.04	31.81 - 32.81	394	396	97
PCB-137	32.07	31.84 - 32.84	360	362	81
PCB-158	32.58	32.00 - 33.00	360	362	81
PCB-138	32.59	32.19 - 33.19	360	362	81
PCB-178	32.42	32.78 - 33.78	394	396	97
PCB-187	33.01	32.85 - 33.85	394	396	97
PCB-183	33.55	33,21 - 34.21	394	396	97
PCB-128	33.88	33.46 - 34.46	360	362	81
PCB-174	34.18	33,76 - 34.76	394	396	97
PC8-177	35.05	34.51 - 36.20	394	396	97
PCB-202	35.41	35.15 - 36.15	430	428	88
PCB-171	35.67	35,50 - 36,50	394	396	97
PCB-156	35.71	35,17 - 36,17	360	362	81
PC8-200	35.78	37.50 - 38,50	430	428	88
PCB-157	36.19	35.90 - 36.90	360	362	81

cont'd...

TABLE 7 - continued

Compound a	RT ^b (min)	RT window	Tgt ^c (m/z)	QI ^d	Abundance ratio ^e
PCBs (native) - cont'd					
PCB-172	36.14	35.84 - 36.84	394	396	97
PCB-180	36.49	36.70 - 37.70	394	396	97
PC8-170/190	36.92	38.64 - 39.64	394	396	97
PCB-201	38.86	38.52 - 39.52	430	428	88
PCB-196/203	39.46	39.16 - 40.16	430	428	88
PCB-208	39.83	39,5 - 40,5	464		
PCB-195	41.74	41.58 - 42.58	430	428	88
PCB-207	41.80	41,35 - 42.35	464		
PCB-194	42.27	41,92 - 42.92	430	428	88
PCB-206	43.14	42.81 - 43.81	464		
PCBs (¹³ C-internal standard)					
¹³ C ₁₂ -PCB-28	18.88	18.58 - 19.58	268		
¹³ C ₁₂ -PCB-52	20.56	20,29 - 21.29	304		
¹³ C ₁₂ -PCB-118	29.19	28.97 - 29.97	338		
¹³ C ₁₂ -PCB-153	30.64	30.43 - 31.43	372		
¹³ C ₁₂ -PCB-180	36,90	36.70 - 37.70	406		
¹³ C ₁₂ ·PCB-194	43,13	42.94 - 43.94	442		
Normalization standard					
¹³ C ₁₂ -PCB-138	31.90	31.40 - 32.40	372		

^a Abbreviations:

PCB - polychlorinated biphenyl

^b Retention time. Oven program: 100°C, hold 3 min; 20°C/min to 180°C; 2.5°C/min to 300°C

^cTarget ion - Mass (m/z)

 $[^]d$ Qualifying ion

^{&#}x27;Abundance ratio - applicable only to ions within molecular ion clusters

7.10. MS Tuning

Note: For detailed instructions on the operation of the instruments, consult the equipment operator's manuals [2.6].

- 7.10.1. Tune the mass spectrometer daily (prior to sample acquisition) with a PFTBA calibration standard (Section 4.7). Detailed procedure and acceptance criteria are described in SOP-CHEM-PROC-12.
- 7.10.2. Print hard copies of the tuning data. An example is given in Figure 2.

7.11. Analysis of Standards and Samples

- 7.11.1. In an autosampler vial, combine 500 μL of the OCs solution prepared in 4.6.3.1 with 50 μL of the ¹³C-labeled chlorobenzenes/PCBs (4.6.1) and 20 μL of the ¹³C₁₂-PCB-138 (4.6.2). Cap the vial and vortex to ensure mixing. *Note:* Prepared just prior to injection used for OCs quantitation.
- 7.11.2. In an autosampler vial, combine 500 μ L of the Aroclor solution prepared in 4.6.3.2 with 50 μ L of the ¹³C-labeled chlorobenzenes /PCBs (4.6.1) and 20 μ L of the ¹³C₁₂-PCB 138 (4.6.2). Cap the vial and vortex to ensure mixing. *Note:* Prepared just prior to injection used for PCBs quantitation.
- 7.11.3. Establish the operating conditions given in Sections 7.7-7.9.
- **7.11.4.** Prepare sequence injection table as described in SOP-CHEM-PROC-10. A typical sequence would consist of the following solutions:
 - OCs quantitation standard (7.11.1)
 - QA reference material
 - Method blank
 - Samples (a series of 10)
 - OCs quantitation standard (7.11.1)
 - PCBs quantitation standard (7.11.2)
 - · QA reference material
 - Method blank
 - Samples (a series of 10)
 - PCBs quantitation standard (7.11.2)

- 7.11.5. Start the injection sequence.
- **7.11.6.** Once the sequence is completed, calculate and print the results as described in SOP-CHEM-DOC-06.

7.12. Initial Calibration Curve

7.12.1. To calibrate the analytical system and determine linearity, inject sequentially, 2 μL of the calibration standard mixtures - CS1, CS2, CS3, CS4 and CS5 (Section 4.6.4), using the instrument conditions listed in 7.7-7.9. An example of calibration curve is given in Figure 3.

Note: Repeat this initial calibration whenever new calibration standard solutions are prepared or if the analytical acceptance criteria for the daily calibration verification standard (CS3) have not been met (Section 10.3).

7.13. Moisture Determination

- 7.13.1. Accurately weigh ca 0.5 g of sample (tissue homogenate) into a preweighed aluminum dish and record the weight to 5 decimal places.
- 7.13.2. Place the dish in a drying oven at 105°C for about two hours, until constant weight is obtained.
- 7.13.3. The calculation of the moisture content is as follows:

% moisture =
$$100 - (W_d/W_w) \times 100$$

where: W_d = weight of dry sample W_w = weight of wet sample

7.14. Lipid Determination - Eggs and tissue samples

Note: Lipid levels are determined to allow calculations of contaminants based on lipid content instead of wet-weight. For plasma samples, see Section 7.15.

- 7.14.1. Allow solvent in the aluminum dish (prepared in Section 7.2.8) to evaporate to dryness in the fume hood.
- 7.14.2. Heat dish in oven at 105°C for 10 min.

- 7.14.3. Take dish out of the oven, allow to cool and reweigh. The difference in weight is the weight of lipids in the sample.
- 7.14.4. The calculation of the lipid content is as follows:

% lipids =
$$(W_1 \times V_1 / W_{te} \times V_{tl}) \times 100$$

where:

 W_1 = weight of lipids (7.14.3)

Wte = weight of sample extracted

Vtl = total volume of extract

 V_1 = volume of extract used for lipid determination (7.2.8)

7.15. Lipid Determination - Plasma samples

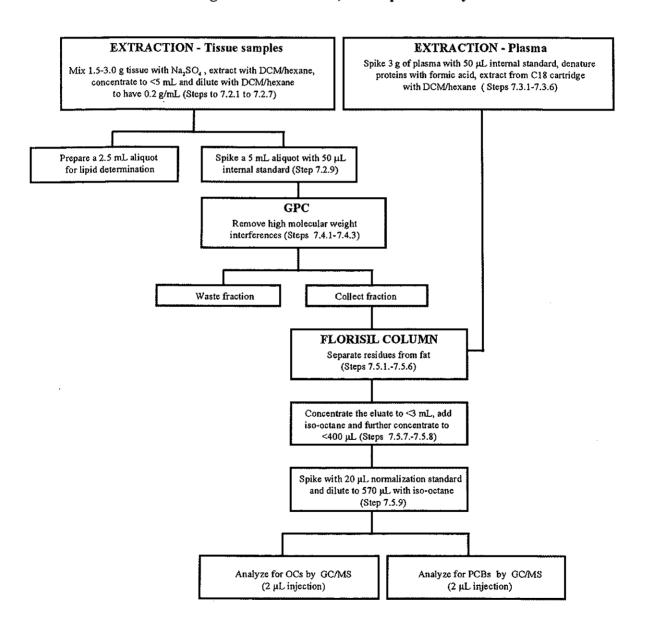
- 7.15.1. Add 20 µL of water (blank), standard olive oil solution of different concentrations (Section 4.8.2) or plasma (unknown) to 15 mL glass centrifuge tubes. Add 0.20 mL of concentrated sulfuric acid to each tube, stopper and mix the contents well on a Vortex mixer.
- 7.15.2. Place all tubes in boiling water bath (Section 5.2.9) for 10 min (±1 min), then cool them in cold water for ca 5 min.
- 7.15.3. Add 10 mL of the phospho-vanillin reagent (Section 4.3.3) to each tube, stopper and mix well on a Vortex mixer. Incubate in a water bath adjusted to 37°C (±2°C) for 15 min.
- 7.15.4. Cool the tubes for ca 5 min and then, within 30 min, measure the absorbance at 540 nm using the following operating parameters:
 - Computer file path: C:\HP8452
 - Mode: F2 Quantitation
 - Wavelength: F1 Single Wavelength (540,0)

Note: Turn the spectrophotometer's lamp on one hour prior to the assay.

- 7.15.5. Press F8 to blank the instrument with the tube containing water.
- 7.15.6. Press F5 (Calibration) to access the "standard information table". Measure the absorbance of each of the standard olive oil solutions to create a calibration curve, specifying the concentration of each one. When readings are done, press F10 to exit.

7.15.7. Press F7 (Analysis) to access the "analysis results table". Measure the absorbance of the unknown plasma samples. The results of the absorbance and concentration will be automatically recorded based on the calibration curve. Print a hard-copy of the results by pressing F9.

FIGURE 1 - Flow diagram of extraction, cleanup and analysis of OCs and PCBs



8. EXPRESSION OF RESULTS

8.1. Calculation of Analyte Concentration

 $C_S = [(A_S / A_{STD}) \times C_{STD}] \times [(V_I / V_F)] \times [(A_{NS \text{ in STD}} / A_{NS \text{ in S}})] \times [(V_{SF} / W_S)] \times 10^{-6}$

where: C_8 = analyte concentration in the sample in $\mu g/g$ (wet weight)

 A_S = area counts of analyte (PCB or OC) in the sample

 A_{STD} = area counts of analyte in the injected standard (7.11.2 for PCBs or 7.11.1 for OCs)

 C_{STD} = analyte concentration in the standard solution (4.6.3.2 for PCBs or 4.6.3.1 for OCs) in $pg/\mu L$

 V_1 = initial volume of the standard solution (4.6.3.2 for PCBs or 4.6.3.1 for OCs) used to prepare the quantitation standard mixture, in μL

V_F = final volume of the PCBs or OCs quantitation standard mixture, in μL

 V_{SF} = final volume of the sample, in μL

 $A_{NS \text{ in STD}}$ = area counts of $^{13}C_{12}$ -PCB-138 normalization standard in the PCBs or OCs quantitation standard mixture

 $A_{NS \text{ in } S}$ = area counts of ${}^{13}C_{12}$ -PCB-138 normalization standard in the sample

W_S = sample weight in g (wet weight)

10⁻⁶ to convert in μg/g

Note: A Microsoft Excel spreadsheet is used to automatically calculate these values following sample acquisition on the GC/MSD.

8.2. Calculation of Recovery for Labeled Internal Standards

The percent recovery (%R) of labeled compounds is calculated as follows:

$$%R = (A_{IS \text{ in S}} / A_{IS \text{ in STD}}) \times (A_{NS \text{ in S}} / A_{NS \text{ in STD}})$$

where: $A_{IS in S}$ = area counts of the labeled internal standard in the sample

 $A_{IS \text{ in STD}}$ = area counts of the labeled internal standard in the quantitation standard mixtures (7.11.1 for OCs and 7.11.2 for PCBs)

 $A_{NS~in~STD}$ = area counts of $^{13}C_{12}$ -PCB-138 normalization standard in the PCBs or OCs quantitation standard mixture

 $A_{NS \text{ in } S}$ = area counts of ${}^{13}C_{12}$ -PCB-138 normalization standard in the sample

Note: Recoveries are automatically calculated by the instrument.

9. REPRESENTATIVE DOCUMENTS

FIGURE 2 - PFTBA spectrum - AutoTune

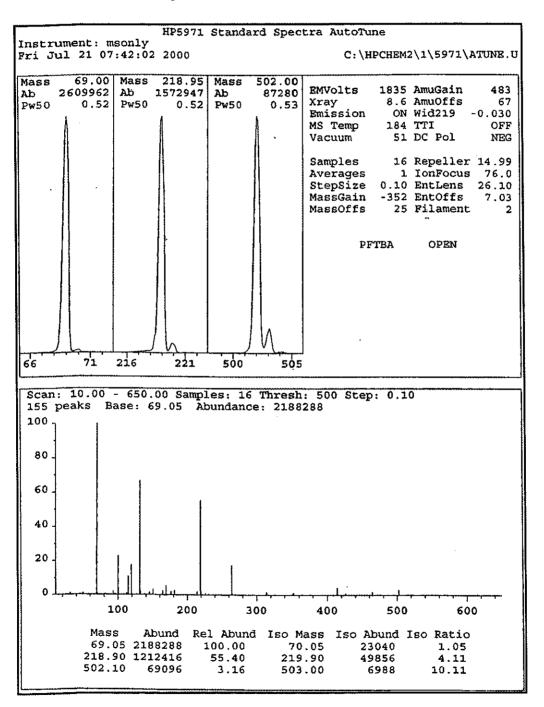
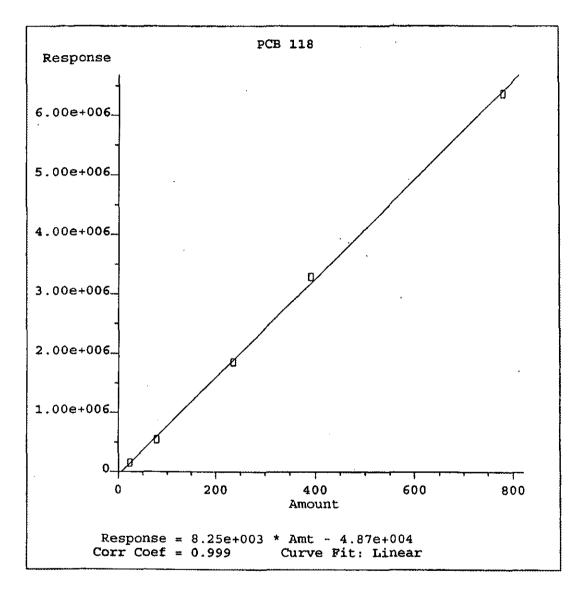


FIGURE 3 - Initial calibration curve for PCB-118



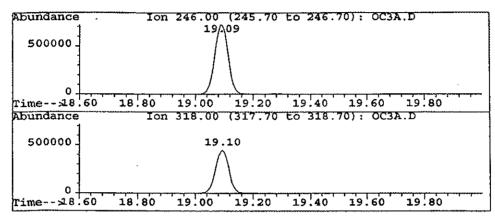
Method Name: C:\HPCHEM2\1\METHODS\5LEVPCB.M Calibration Table Last Updated: Mon Jul 24 14:46:12 2000

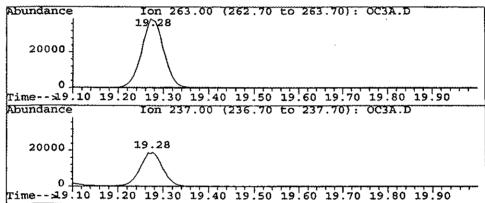
Note: Actual concentrations shown in Table 3.

FIGURE 4 - Chromatogram of OCs standards

Name of file: C:\HPCHEM2\1\DATA\LINEAR\OC3A.D

Operator: H.W. Sample: OC3A.D





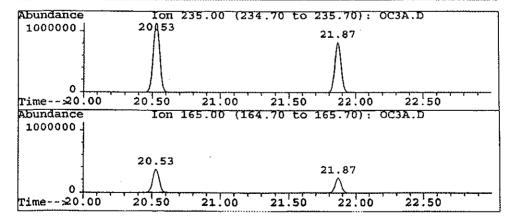
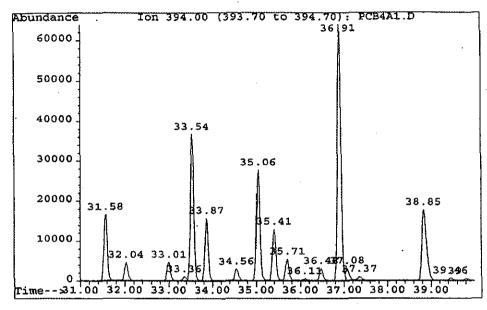


FIGURE 5 - Chromatogram of PCBs standards (Aroclor mixture)

Name of file: C:\HPCHEM2\1\DATA\LINEAR\PCB4A1.D

Operator: H.W. Sample: PCB4A1.D



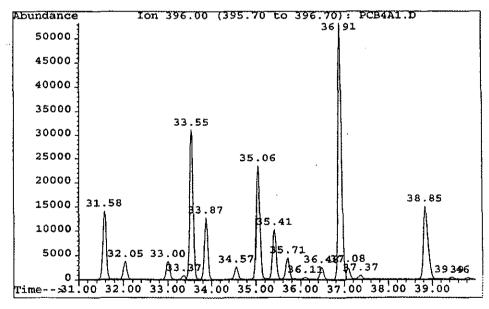
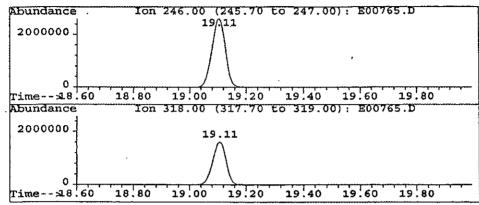
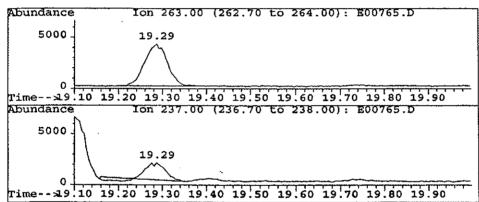


FIGURE 6 - Chromatogram of herring gull egg sample (analyzed for OCs)

Name of file: C:\HPCHEM2\1\DATA\0016A\E00765.D

Operator: H.W. Sample: E00765.D





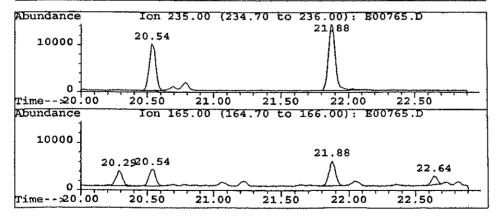
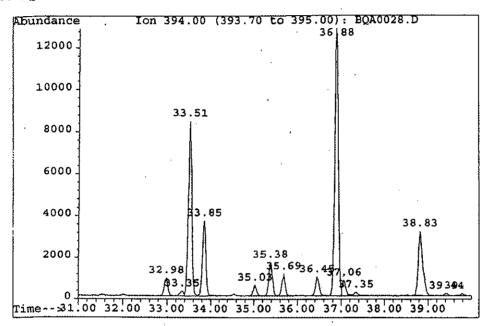
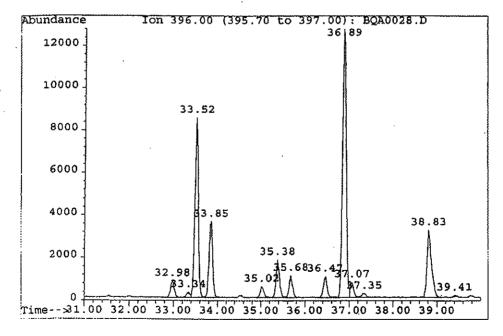


FIGURE 7 - Chromatogram of herring gull egg sample (analyzed for PCBs)

Name of file: C:\HPCHEM2\1\DATA\0028\BQA0028.D

Operator: H.W. Sample: BQA0028.D





10. QUALITY CONTROL

10.1. MSD Tuning

The MSD is tuned weekly with the PFTBA calibration standard using the AutoTune program, and daily with the QuickTune program. The tuning of the instrument must meet the criteria for conformance outlined in SOP-CHEM-PROC-12 before sample analysis. Tune files are archived in a logbook.

10.2. Calibration Verification

A five-point initial calibration standard curve is made with the OCs and PCBs standard mixtures to cover the range of interest. This established calibration curve is verified daily, by analyzing a calibration verification standard (quantitation standard) having a mid-point concentration. The calculated concentration of each compound must be within 20% of its actual known value. The final concentration of any reportable compounds must be within the demonstrated linearity of the detector. If necessary, samples are diluted with iso-octane to meet the calibration range.

10.3. Detection Limits and Reporting Limits

A nominal or minimum detectable concentration (MDC) is usually described as the concentration of analyte which produces a signal in an instrument 3 times the average noise level. In this multi-residue method, it is possible but not practical to list the detection limits for each compound of interest. Variability between compounds arises due to varying background noise and response factors for each compound due to the different mass ions being monitored. The background noise is affected by several factors, such as tissue size and tissue type, and instrument effects such as column bleed and cleanliness of the source on the day of the analysis.

As a general rule, a detection limit of at least 0.001 PPM is achievable for all compounds. For the purposes of reporting data, no results less than this concentration are reported and a result of ND (not detected) appears in the Laboratory Services Section analytical test report. If a computed result falls in the range 0.0001 and 0.0009 PPM, the compound is defined as being detected but the result would be too variable to be reliable so a designation of TR (trace) is listed beside the compound in the final report.

10.4. Ongoing Precision and Recovery

An aliquot of the QA Reference Material (Herring gull eggs - Section 4.9) is analyzed along with each batch of samples. The concentration of the major compounds (PCB-52, PCB-66, PCB-101, PCB-110, PCB-149, PCB-118, PCB-146, PCB-153, PCB-138, PCB-187, PCB-180, PCB-170, PCB-201, PCB-203, HCB, p,p'-DDE, photo-mirex, mirex, oxychlordane, cis-nonachlor, heptachlor epoxide and dieldrin) is determined and the results are compared to the previously established acceptance limits (i.e., \pm 2 SD of the long-term mean plotted in a Shewart chart - ref. SOP-CHEM-DOC-02).

To determine the degree of analyte loss during sample cleanup, each sample (including the PCBs and OCs standard mixtures 7.11.1 and 7.11.2) is spiked with ¹³C-labeled chlorobenzenes/PCBs internal standard mixture (4.6.1). It is assumed that during sample cleanup, native compounds behave the same way as the labeled compounds. Analysis is accepted when the % internal standard recoveries for most PCBs and OCs are between 80% and 110%, and for the highly volatile compounds (eluting in the 1st, 2nd and 3rd chromatographic windows) are over 60%. The levels of contaminants reported in the analytical test report are not corrected for internal standard recoveries, but the recovery data are included.

10.5. Accuracy

To improve the accuracy of the method, the compound levels can be corrected based on the recovery of the ¹³C-labeled chlorobenzenes/PCBs internal standard mixture (4.6.1). The majority of the chlorinated pesticides and PCB congeners levels can be corrected based on the average recovery of the five higher chlorinated ¹³C-labeled PCB standards (i.e., PCB-52, PCB-118, PCB-153, PCB-180 and PCB-194). The more volatile compounds can be recovery-corrected using the recovery of their corresponding ¹³C-labeled standards. For volatile compounds without ¹³C-surrogates, retention time interpolation of recoveries with the ¹³C-standards can be used.

The accuracy of the quantitation standards is verified annually with a second source standard (containing most of the congeners of interest) as described in SOP-CHEM-PROC-13.

10.6. Method Blank

A method blank is run with each batch of samples to determine the levels of contamination associated with the processing and analysis of samples. If

problems with blank exist, associated data are carefully evaluated and appropriate corrective actions are applied. Blank values are not subtracted from reportable values. A compound found in a blank and also in an associated sample is flagged in the analytical test report when present at a ratio of at least 5/1, sample to blank.

10.7. Target Compound Identification

Peak GC retention time must be within their pre-defined retention time windows.

All ions present in the standard mass spectrum at a relative intensity greater than 10% must be present in the sample spectrum.

The relative intensities of these ions must agree within $\pm 20\%$ between the standard and sample spectra (e.g., for an ion with an abundance of 50% in the standard spectrum, the corresponding sample ion abundance must be between 30% and 70%).

10.8. Standard Operating Procedures

Other related SOPs relevant to this analytical method:

- SOP-CHEM-DOC-01: for the use of log-books
- SOP-CHEM-DOC-02: for the creation of control charts for quality control samples
- ◆ SOP-CHEM-DOC-04: for the archival of gas chromatography data files
- SOP-CHEM-DOC-05: for the archival of analytical test reports
- SOP-CHEM-PROC-01: for pipets calibration
- SOP-CHEM-PROC-03: for balances calibration
- SOP-CHEM-PROC-05 for the preparation and storage of standard solutions
- SOP-CHEM-PROC-06: for the monitoring of refrigerator's temperatures
- SOP-CHEM-PROC-07: for glassware cleaning
- SOP-CHEM-PROC-10: for running a sequence with the GC/MSD
- SOP-CHEM-PROC-12: for the tuning and calibration of the MSD
- SOP-CHEM-PROC-13: for verification of standard with a second source standard
- SOP-CHEM-MAIN-04: for the maintenance of the GC/MSD

10.9. Data Validation

Data validation is ensured by an internal quality assurance audit done by an independent reviewer (Head of the Laboratory Services Section), before the release of the analytical test report. Results of this verification are recorded on the "Data Validation Form for OC/PCBs Reports".

If large discrepancies in the analytical data between the specimens from close geographical areas are noted, then the raw data are examined - re-analysis of the sample aliquot may be indicated.

10.10.Method Validation

Some method validation data related to linearity and detection limits are compiled in internal reports [2.3 and 2.4]. Linear curves were obtained with OCs mixed standards solutions ranging from ca 10 to 1000 pg/ μ L. The limits of detection varied, depending on the compound, but ranged from a low of 10 pg/ μ L to 60 pg/ μ L for a 2 μ L injection. For the PCBs, mixed Aroclor standards solutions ranging from 300 to 30 000 pg/ μ L (for the sum of the congeners), were analyzed. Most congeners responded linearly (R² between 0.991 and 1.00). The limits of detection varied depending on the congener but ranged from 1 pg for dichlorobiphenyls to 20 pg for nonachlorobiphenyl (PCB-206) for a 2 μ L injection.

Recoveries experiments using spiked chicken eggs are detailed in Reference 2.5 - Internal report CHEM-OC-97-34.

10.11. External Quality Control

Whenever appropriate inter-laboratory check sample programs are available, the Laboratory participates in them. One example is "The Analysis of PCB Congeners and Organochlorines in Biota", which includes a variety of fish tissues provided as part of the Northern Contaminants Program (Indian and Northern Affairs Canada) [2.10].

11. CRITICAL CONTROL POINTS

A number of critical points are identified at various steps in Section 7. The following are repeated for emphasis:

Sample extracts must not be allowed to evaporate to dryness at anytime during the sample cleanup procedure. Several of the residue compounds, such as the chlorobenzenes and low chlorinated PCBs, are volatile. Reducing extracts to dryness will therefore result in loss of analytes. Evaporation of sample extracts using directed gas streams should be avoided for the same reasons, especially at volumes less than 1 mL. It is always preferable to use a rotary evaporator with a suitable flask volume, or sequence of flasks.

Sufficient mixing of the deactivated Florisil is necessary in order to assure equilibrium, before using the Florisil for sample cleanup.

To maintain the consistency and reproducibility of the GPC cleanup, always use the EnvirobeadsTM S-X3 described in 4.2.1.

Trace contaminant levels are determined by this method and the elimination of interferences is essential. They could occur through sample handling, reagents, solvents, instruments or labware.

MULTIRESIDUE METHOD FOR THE DETERMINATION OF CHLORINATED PESTICIDES AND POLYCHLORINATED BIPHENYLS (PCBS) IN POLAR BEAR TISSUES

1. SCOPE AND FIELD OF APPLICATION

The method that follows has been used in our laboratory for the analysis of various polar bear tissues such as fat, liver, muscle, lung, kidney, brain, gonads and milk for the chlorinated pesticides and polychlorinated biphenyls (PCBs) that are known to be present in the Canadian wildlife [2.4, 2.5]. Four different extraction methods are described to accommodate tissues with various lipids content, sample size and matrix. The specific contaminants targeted are listed in **Table 2**. Typical detection limits are in the range of 0.005 to 0.01 ppm.

2. REFERENCES

- 2.1. Zhu, J.P., Simon, M., Mulvihill, M.J. and Norstrom, R.J. (1994) Single fraction determination of PCBs and organochlorine contaminants by GC-MS using characterized tissue extracts as secondary standards. NWRC unpublished data.
- 2.2. Won, H. (1994) Evaluation of GC/MSD method for 21 organo chlorine standards for detection limit and linear range. NWRC internal report.
- 2.3. Won, H. (1994) Evaluation of GC/MSD method for some isomers in Aroclor (1242/1254/1260) (1:1:1) for detection limit and linear range. NWRC internal report.
- 2.4. Muir, D.C.G., Norstrom, R.J. and Simon, M. (1988) Organochlorine contaminants in Arctic marine food chains: Accumulation of specific PCB congeners and chlordane-related compounds. *Environ. Sci. Technol.*, 22, 1071-1079.
- 2.5. Norstrom, R.J., Simon, M., Muir, D.C.G. and Schweinsburg, R. (1988) Organochlorine contaminants in Arctic marine food chains: Identification, geographical distribution and temporal trends in polar bears. *Environ. Sci. Technol.* 22, 1063-1071.
- 2.6. Norstrom, R.J., Belikov, S.E., Born, E.W., Garner, G.W., Malone, B., Olpinski, S., Ramsay, M.A., Schliebe, S., Stirling, I., Stishov, M.S., Taylor, M.K., Wiig, Ø. (1998) Chlorinated hydrocarbon contaminants in polar bears from eastern Russia, North America, Greenland, and Svalbard: Biomonitoring of Arctic pollution. Arch. Environ. Contam. Toxicol. 35, 354-367.
- 2.7. HP 5970B Mass selective detector hardware manual (05970-90049) and MS ChemStation user's guide (HP-G1034-90043).

- 2.8. O.I. Analytical (1992) Operating manual, Autoprep 1000TM chromatograph, Rev. 2.
- **2.9.** Zhu, J.P. (1993) Comparability of analytical data for PCBs and chlorinated pesticides: Review of polar bear fat extract results. NWRC internal report.

3. PRINCIPLES AND DEFINITIONS

Tissues are dried out with sodium sulfate and the residues are extracted with DCM/hexane. Milk samples are first denatured with methanol, and then extracted with DCM/hexane. Removal of lipids and biogenic compounds from the lipid extracts is done by gel permeation chromatography (GPC). The eluant is further cleaned up by Florisil column chromatography. Finally, the OCs and PCBs are determined via high resolution gas chromatography coupled to a mass spectrometry detection system. Identification of target analytes is accomplished by comparing GC retention times and specific mass fragments known to be present in the spectra of authentic compounds. Quantitation is accomplished by comparing the intensity of targeted mass fragments in specimen extracts to the same compounds in standard mixtures or in polar bear reference material (Section 4.5) previously characterized at NWRC [2.1].

4. REAGENTS, SOLUTIONS AND STANDARDS

SAFETY PRECAUTIONS

- ⇒ The toxicity or carcinogenicity of each reagent and standard used in this method has not been precisely defined. Each chemical must be treated as a potential health hazard and exposure should be reduced to the lowest possible level.
- ⇒ Operations with toluene, DCM, hexane, methanol and acetone should be performed in a fume hood and dermal contact with solvents should be avoided.
- ⇒ Standards should always be opened and used in a fume hood. Handling of these compounds must be done only by qualified technical staff.
- ⇒ General safety rules and waste disposal procedures that apply to the Trace Organic Chemistry Laboratory must be followed (ref. Safety Manual).
- ⇒ Material Safety Data Sheets (MSDSs) for the products used in the assay must be read.

4.1. Reagents

- 4.1.1. Acetone, Omnisolv®, EM SCIENCE CAAX0116-1
- 4.1.2. Hexane, Omnisolv®, EM SCIENCE CAHX0296-1
- 4.1.3. Dichloromethane, Omnisolv®, EM SCIENCE CADX0831-1
- 4.1.4. Methanol, Omnisolv®, EM SCIENCE CAMX0488-1
- 4.1.5. Toluene, Omnisolv®, EM SCIENCE CATX0737-1
- 4.1.6. Diethyl ether, anhydrous, Omnisolv®, EM SCIENCE CAACS291-76
- **4.1.7.** 2,2,4-Trimethylpentane (iso-octane), Omnisolv[®], EM SCIENCE CATX1389-1
- **4.1.8.** Sodium sulfate, anhydrous granular, (Na₂SO₄), EM SCIENCE ACS850-46

Wash 600 g of Na_2SO_4 in a glass column 3 cm ID x 50 cm long with 600 mL DCM/hexane (1:1), air dry in an open dish under the fume hood, heat 3 h at 400°C in a muffle furnace, cool and transfer in a tightly capped glass bottle. *Note:* If, after heating, the sodium sulfate develops a greyish cast (due to the presence of carbon in the crystal matrix), discard that batch.

- **4.1.9.** De-ionized water from the Milli-RO / Milli-O system (Millipore)
- 4.1.10. Helium, compressed bottled gas, Praxair, HE UHP SG 103168K
- 4.1.11. Nitrogen, compressed bottled gas, Praxair, N₂ PRE PURE SG 105411K

4.2. Adsorbents for Sample Cleanup

See Section 4.2 of MET-CHEM-OC-04.

4.3. Solutions

- **4.3.1.** DCM/hexane, 1:1 (v/v)
- **4.3.2.** Potassium oxalate 0.6 M Dissolve 55.2 g in de-ionized water and dilute to 500 mL.

4.4. Standards

Internal standard spiking solution and normalization standard as described in Sections 4.6.1 and 4.6.2 of MET-CHEM-OC-04.

4.5. QA Reference Material - Secondary Quantitation Standard

Since certified reference material with the same matrix does not exist, adipose tissue from polar bear prepared in-house is used. The current reference material used, identified as PBQ2, is from polar bear PB-142 collected in 1984 in the Canadian Arctic. Details about preparation and quantitation of the extracts is described in Norstrom et al. [2.6].

4.6. Method Blank

See Section 4.10 of MET-CHEM-OC-04.

5. AUXILIARY EQUIPMENT

5.1. Glassware and Labware

As described in Sections 5.1 of MET-CHEM-OC-04 with the following exceptions:

- a) Item 5.1.4 (C18 cartridges) are not used
- b) Filter disks 0.2 μm (13 mm diam.), attached to a 5 mL syringe, are needed for adipose tissue biopsy samples (ref. Section 7.4.8).

5.2. Equipment

As described in Sections 5.2 of MET-CHEM-OC-04 with the following exceptions:

- a) Items 5.2.8-5.2.10 (Visiprep solid phase extraction vacuum and water baths at 100°C and 30°C) are not used.
- b) Omni-mixer homogenizer with 400-mL cup Diamed (Sorvall 17105) is used for large adipose tissue samples (Section 7.3.1).

5.3. Instrumentation

See Sections 5.3.1-5.3.2 of MET-CHEM-OC-04.

6. SPECIMEN OR SAMPLE HANDLING REQUIREMENTS

Samples are prepared as described in the Tissue Preparation Unit's standard operating procedure SOP-TP-PROC-07. Collection and preservation protocols are described in Norstrom et al [2.6] and in "Protocol for Field Collection and Storage of Wild Birds for Biomarker Studies" (S. Trudeau, Biomarker Laboratory, NWRC, 1992).

7. PROCEDURE

7.1. Columns Preparation

See Sections 7.1.1 and 7.5.1-7.5.3 of MET-CHEM-OC-04.

7.2. Extraction - Milk samples

- 7.2.1. Accurately weigh 3.00 g of milk in a 50 mL screw-top centrifuge tube and spike with 50 μ L of the ¹³C-labeled chlorobenzene/PCB internal standard spiking solution.
- 7.2.2. Add 3 mL of the $K_2C_2O_4$ solution (Section 4.3.2) to the spiked milk sample and mix with Vortex mixer.
- 7.2.3. Add 3 mL of methanol to the mixture, and let it stand for 15 min. Mix thoroughly with Vortex and let it stand for another 30 min.
- 7.2.4. Add 6 mL diethyl ether and 6 mL hexane. Vortex the mixture.
- 7.2.5. Centrifuge for 2 min at ca 2000 RPM to separate the organic and the aqueous phases.
- 7.2.6. Transfer the organic layer (top layer) with a Pasteur pipet into a 125 mL round bottom flask. Repeat the extraction procedure (steps 7.2.4-7.2.5) two more times.
- 7.2.7. Add 15 mL hexane and ca 1 g of anhydrous Na₂SO₄ (Section 4.1.8) to the combined extracts in order to absorb any residual moisture. Let

stand for 15 min.

- 7.2.8. Concentrate the extract to less than 5 mL using the rotary evaporator with the water bath adjusted to ca 30°C.
- **7.2.9.** Quantitatively transfer the extract into a graduated 15 mL centrifuge tube and dilute to 10 mL with DCM/hexane solution (1:1).
- 7.2.10. Transfer a 1 mL aliquot (equivalent to 0.3 g sample) in a pre-weighed aluminum dish, to be used for lipid determination as described in Section 7.14 of MET-CHEM-OC-04.
- 7.2.11. The rest of the extract is ready for GPC cleanup. *Note*: since 10% of the sample is removed for lipid determination, the internal standard % recovery measured will have to be corrected (see Section 10.4).
- 7.3. Extraction Adipose tissue samples (larger than 5 g wet weight)
 - 7.3.1. Accurately weigh ca 5 g of sample (still frozen) in a 400 mL Sorval cup. Add 50 mL DCM/hexane (1:1) and homogenize with Omni mixer for ca 10 min.
 - 7.3.2. Plug the end of a powder funnel with some treated glass wool (Section 5.1.12) and add ca 25 g of Na₂SO₄.
 - 7.3.3. Place a 250 mL round bottom evaporating flask under the funnel.
 - 7.3.4. Decant the homogenized sample mixture onto the Na₂SO₄, keeping the connective tissue in the Sorval cup.
 - 7.3.5. Add another 50 mL DCM/hexane (1:1) in the cup and homogenize for ca 10 min. Decant the solvents onto the Na₂SO₄.
 - 7.3.6. Repeat step 7.3.5 and transfer solvents and connective tissue on top of Na₂SO₄ layer.
 - 7.3.7. Rinse funnel with 20 mL DCM/hexane (1:1).
 - 7.3.8. Evaporate the eluate to less than 5 mL on a rotary evaporator with water bath at ca 30°C, then quantitatively transfer into a 12 mL graduated centrifuge tube. Rinse the flask three times with ca 1 mL of DCM/hexane (1:1), adding the rinses to the centrifuge tube.

- 7.3.9. Adjust volume to 10 mL with DCM/hexane (1:1) to have a sample concentration of 0.5 g/mL.
- 7.3.10. Transfer a 1 mL aliquot (equivalent to 0.5 g tissue) in a pre-weighed aluminum dish, to be used for lipid determination (ref. Section 7.14 of MET-CHEM-OC-04).
- 7.3.11. Pipette a 2 mL aliquot into a GPC tube. Spike the extract (equivalent to 1 g tissue wet weight) with 50 µL of the ¹³C-labeled chlorobenzene/ PCB internal standard spiking solution, and dilute to 10 mL with DCM/hexane (1:1). The sample is now ready for GPC cleanup.
- **7.3.12.** Transfer the remaining extract from 7.3.9 in a 10 mL sealed ampule for future use.
- 7.4. Extraction Adipose tissue samples (biopsy samples)
 - 7.4.1. Accurately weigh ca 100 mg (or less) of fat, previously detached from the surface skin and hair, into a pre-weighed 20 mL scintillation vial. *Note:* Separate skin from the fat with a sharp scalpel while the biopsy sample is still partially frozen to facilitate the task.
 - 7.4.2. Add ca 1 g of the treated anhydrous Na₂SO₄ and re-weigh the vial.
 - **7.4.3.** Grind the mixture with a stainless steel spatula. Cover vial with aluminum foil (leaving spatula in the vial) and allow to stand for 30 min.
 - 7.4.4. Spike the mixture with 20 μ L of the ¹³C-labeled chlorobenzene/PCB internal standard spiking solution.
 - 7.4.5. Add 5 mL of DCM/hexane (1:1) and allow to stand for 30 min. Transfer extracting solvents with a Pasteur pipet in into a 125 mL round bottom flask.
 - 7.4.6. Repeat extraction (Step 7.4.5) three more times, combining the extracts in the 125 mL flask. *Note:* Thoroughly mix after each addition of fresh solvents to ensure complete lipid extraction.
 - 7.4.7. Concentrate the extract to less than 2 mL using the rotary evaporator with the water bath adjusted to ca 30°C.
 - 7.4.8. Filter the concentrated extract through a 0.2 µm filter disk attached to a

5 mL syringe. Rinse the filter 3 times with 1 mL DCM/hexane (1:1), combining all fractions in a GPC tube. The sample is ready for GPC cleanup.

7.5. Extraction - Low-lipid samples

Note: This extraction method is used for low-lipid samples such as liver, muscle, kidney, gonad, brain and lung.

See Sections 7.2 of MET-CHEM-OC-04

7.6. Sample Cleanup by GPC

Spiked sample extract (from 7.2.11, 7.3.11, 7.4.8 or 7.5) is injected into one loop of the GPC as described in Section 7.4 of MET-CHEM-OC-04.

7.7. Sample Cleanup by Florisil Column

As described in Sections 7.5.1-7.5.4 of MET-CHEM-OC-04 and then:

- 7.7.1. Load extract obtained after GPC cleanup step on top of the Florisil column using Pasteur pipet. Rinse the flask 3-4 times with small portions of DCM/hexane (1:1). Add rinses to column, allowing solvent to drain to packing level in between rinses.
- 7.7.2. As last rinse approaches top of adsorbent layer, add 70 mL of DCM/hexane (1:1) and allow elution to proceed at ca 5 mL/min.
- 7.7.3. Concentrate the eluate to less than 2 mL with rotary evaporator and quantitatively transfer to a pear-shaped flask and concentrate to less than 0.5 mL with rotary evaporator. Quantitatively transfer in a 1 mL reacti-vial, rinsing the flask with iso-octane.
- 7.7.4. Further concentrate the eluate from the milk, low-lipid and small biopsies samples to less than 100 µL under a gentle nitrogen flow and to less than 200 µL for the large adipose tissues.
- 7.7.5. Quantitatively transfer the sample (with a Pasteur pipet) to a 500 μ L Reacti-vial previously marked at the 270 μ L level for the large adipose tissue samples and to the 170 μ L for the others. Rinse flask with a few drops of iso-octane, adding the rinses to the vial.

- 7.7.6. Add exactly 20 μL ¹³C₁₂-PCB-138 normalization standard to the Reactivial using an Eppendorf pipet. Adjust the final volume to the 170 or 270 μL mark with iso-octane. *Note:* Exact volume is not critical, since normalization standard is used to quantify residue levels.
- 7.7.7. Cap and vortex until thoroughly mixed. The sample is ready to inject into the GC/MS (see Section 7.13.3).

7.8. GC Operating Conditions

7.8.1. Column

See Section 7.6.1 of MET-CHEM-OC-04.

7.8.2. Injection information

See Section 7.6.2 of MET-CHEM-OC-04.

- 7.8.3. Oven temperature programme
 - 100°C, hold 3 min; 20°C/min to 180°C; 5°C/min to 300°C
- 7.8.4. Carrier gas (He)

See Section 7.6.4 of MET-CHEM-OC-04

7.9. MSD Operating Conditions

See Section 7.7.1 of MET-CHEM-OC-04

7.10. Data Analysis Parameters

Calibration settings
 Reference window: 10,00%

Non-reference window
 5.00%

• Default multiplier 1.00

• Default sample concentration 1.00

7.11. Compound Information

Chromatographic windows and characteristic ions are given in **Table 1**. Retention time, target ion and qualifying ion are given in **Table 2**. The list is based on those compounds actually found in polar bear tissues using PBQ2 as a secondary standard.

Table 1. Chromatographic windows and characteristic ions

SIM Group	Start time (min)	Dwell time (ms)	lons
Group 1	7.40	60	216, 224 ^a
Group 2	9.40	60	250, 258 ^a
Group 3	10.20	60	183, 219, 250, 284, 292 a
Group 4	13.00	60	238, 268 ^a , 292, 304 ^a
Group 5	16.00	60	255, 291, 292, 308, 339, 353, 378, 387
Group 6	17.60	60	292, 326, 339, 373, 375
Group 7	18.40	60	246., 326, 373, 375, 409
Group 8	19.00	60	246, 326, 360, 380
Group 9	20.10	60	235, 237, 326, 338 ^a , 360, 443,
Group 10	20.70	60	235, 237, 326, 360, 372 ^a , 394,
Group 11	21.70	60	235, 237, 360, 394,
Group 12	23.50	60	360, 394, 406 ^a
Group 13	25.80	60	251, 311, 430, 442 ^a , 464, 498

^aions selected to monitor the ¹³C-labeled internal standards

TABLE 2 - Retention time and target ions for secondary quantitative standard (PBQ2) and internal standard

Compound ^a	RT ^b (min)	RT window	Tgt ^c (m/z)	QI ^d	Abundance ratio ^e
Quantitation standard (PBQ2)					
1,2,4,5-tetrachlorobenzene	7.71	7.31 - 8.11	216	214	77
pentachlorobenzene	9.75	9.35 - 10.15	250	252	65
hexachlorobenzene	12.15	11.75 - 12.65	284	286	81
aipha-hexachlorocyclohexane	11.9	11.5 - 12.3	219	183	80
beta-hexachlorocyclohexane	12.55	12.15 - 12.85	219	183	119
octachlorostyrene	17.11	16.71 - 17.51	308	380	42
compound C	14.31	13.91 - 14.71	238	373	
U-3	10.47	10.07 - 10.87	250	252	85
photoheptachtor	16.62	16.22 - 17.02	339		
heptachlor epoxide	17.27	16.87 - 17.67	353	237	
oxychlordane	17.32	16.92 - 17.72	387	119	
U-4	17.48	17.08 - 17.88	255	291	34
C-5	17.87	17.62 - 18.12	373	375	96
C-3	17.89	17.49 - 18.29	339		
C-4	18.25	17.85 - 18.65	373	375	89
nonachlor III	18.57	18.27 - 18.87	409	407	88
t-nonachior	18.75	18.45 - 19.05	409	407	88
U-2	20.39	19.99 - 20.79	443		
dieldrin	19.39	18.99 - 19.79	380	308	
p,p'-DDE	19.21	18.81 - 19.61	246	318	95
p,p'-DDD	20.65	20.25 - 21.05	235	237	42
p,p'-DDT	21,99	21.59 - 22.39	235	237	42
PCB-47/48	15.64	15.24 - 16.04	292	290	77
PCB-74	17.22	16.82 - 17.62	292	290	77
PCB-56/60	18.08	17.68 - 18.48	292	290	77
PCB-99	18.43	18.23 - 18.83	326	328	65
PCB-85	19.32	18.92 - 19.72	326	328	65
PCB-154	19.53	19.13 - 19.93	360	362	81
PCB-149	20.35	19.95 - 20.75	360	362	81
PCB-118	20.40	20.00 - 20.8	326	328	65
PCB-146	20.97	20.57 - 21.37	360	362	81
PCB-163	21.19	20.79 - 21.59	360	362	81
PCB-105	21.34	20.94 - 21.49	326	328	65
PCB-137	21.89	21,49 - 22.29	360	362	81
PCB-138/163	22.16	21,76 - 22.56	360	362	81
PCB-187	22.76	22.36 - 23.16	394	396	97
PCB-183	22,94	22.54 - 23.34	394	396	97
PCB-156	23.97	23,57 - 24.37	360	362	81
PCB-157	24.17	23.77 - 24.57	360	362	81
PCB-180	24.55	24.15 - 24.95	394	396	97
PCB-170/190	25.6	25.20 - 26.00	394	396	97
PCB-195	27.16	26.66 - 27.76	430	428	88
PCB-194	27.82	27.42 - 28.22	430	428	88
PCB-206	29.14	28.74 - 29.54	464		
PCB-209	30.20	29.60 - 30.80	498		1

cont'd

TABLE 2 - continued

Compound ^a	RT ^b (min)	RT window	Tgt ^c (m/z)	QI ^d	Abundance ratio ^e
Internal standard					
¹³ C ₆ -1,2,4,5-tetrachlorobenzene	7.71	7.31 - 8.11	224		
¹³ C ₆ -1,2,4,5-pentachlorobenzene	9.75	9.35 - 10.15	258		
¹³ C ₆ -1,2,4,5-hexachlorobenzene	12.15	11.75 - 12,55	292		
¹³ C ₁₂ -PCB-28	14.35	13.95 - 14.75	268		
¹³ C ₁₂ -PCB-52	15.39	14.99 - 15.79	304		
¹³ C ₁₂ -PCB-118	20.4	20.00 - 20.80	338		
¹³ C ₁₂ -PCB-153	21.18	20.78 - 21,58	372		
¹³ C ₁₂ -PCB-180	24.54	24.14 - 24.94	406		
¹³ C ₁₂ -PCB-194	27.82	27.42 - 28.22	442		

^a Abbreviations:

compound C - unknown structure

U2, U3, U4 - unknown chlorinated compounds

C3,C4, C5 - unknown chlordane related compounds

DDE - dichloro diphenyl dichloro ethylene

DDD - dichloro diphenyl dichloro ethane

DDT - dichloro diphenyl trichloro ethane

PCB - polychlorinated biphenyl

^b Retention time. Temperature program: 100°C, hold 3 min; 20°C/min to 180°C; 5°C/min to 300°C

Target ion - Mass (m/z)

^dQualifying ion

^{&#}x27;Abundance ratio - applicable only to ions within molecular ion clusters

7.12. MS Tuning

See Section 7.10 of MET-CHEM-OC-04.

7.13. Analysis of Standards and Samples

- 7.13.1. Quantitative standard for milk samples, adipose tissues from biopsy and low-lipid samples In a 200 μL insert of an autosampler vial, combine 140 μL of the quantitative standard PBQ with 20 μL of the ¹³C-labeled chlorobenzenes/PCBs and 10 μL of the normalization standard ¹³C₁₂-PCB-138, for a final volume of 170 μL. Cap the vial and vortex to ensure mixing. *Note:* Prepared just prior to injection.
- 7.13.2. Quantitative standard for larger adipose tissue sample In a 300 μL insert of an autosampler vial, combine 240 μL of the quantitative standard PBQ with 20 μL of the ¹³C-labeled chlorobenzenes/PCBs and 10 μL of the normalization standard ¹³C₁₂-PCB-138, for a final volume of 270 μL. Cap the vial and vortex to ensure mixing. Note: Prepared just prior to injection.
- 7.13.3. Samples Transfer the spiked sample from Section 7.7.7 into a 200 (or 300) μL insert of an autosampler vial and cap. Store in the dark at room temperature until ready for analysis. If analysis will not be performed on the same day, store the vial at 0-4°C.
- 7.13.4. Establish the operating conditions given in Sections 7.8-7.9.
- 7.13.5. Prepare sequence injection table as described in SOP-CHEM-PROC-10. A typical sequence would consist of the following solutions:
 - PBQ quantitation standard (7.13.1 or 7.13.2)
 - · Method blank
 - Samples (a series of 5)
 - PBQ quantitation standard (7.13.1 or 7.13.2)

8. EXPRESSION OF RESULTS

8.1. Calculation of Analyte Concentration

 $C_S = [(A_S / A_{PBQ}) \times C_{PBQ}] \times [(V_{PBQI} / V_{PBQF})] \times [(A_{NS \text{ in } PBQ} / A_{NS \text{ in } S})] \times [(V_{SF} / W_S)] / 10^{-6}$

where: C_S = analyte concentration in the sample in $\mu g/g$ (wet weight)

 A_S = area counts of analyte in the sample

A_{PBQ} = area counts of analyte in the PBQ injected standard

 C_{PBQ} = analyte concentration in the PBQ standard solution in pg/ μ L

 V_{PBQI} = initial volume of the PBQ standard solution used to prepare the quantitation standard mixture, in μL

 V_{PBOF} = final volume of the PBQ quantitation standard mixture, in μL

 V_{SF} = final volume of the sample, in μL

 $A_{\rm NS~in~PBQ}$ = area counts of $^{13}C_{12}$ -PCB-138 normalization standard in the PBQ standard solution

 $A_{NS \text{ in } S}$ = area counts of ${}^{13}C_{12}$ -PCB-138 normalization standard in the sample

Ws = sample weight in g (wet weight)

10⁻⁶ to convert in µg/g

Note: A Microsoft Excel spreadsheet is used to automatically calculate these values following sample acquisition on the GC/MSD.

8.2. Calculation of Recovery for Labeled Internal Standards

The percent recovery (%R) of labeled compounds is calculated as follows:

$$%R = (A_{IS \text{ in } S} / A_{IS \text{ in } PBO}) \times (A_{NS \text{ in } S} / A_{NS \text{ in } PBO})$$

where: $A_{IS \text{ in } S}$ = area counts of the labeled internal standard in the sample

 $A_{IS\ in\ PBQ}$ = area counts of the labeled internal standard in the quantitation standard mixture

 $A_{NS \text{ in } S}$ = area counts of ${}^{13}C_{12}$ -PCB-138 normalization standard in the sample

 $A_{NS \text{ in PBQ}}$ = area counts of $^{13}C_{12}$ -PCB-138 normalization standard in the quantitation standard mixture

Note: Recoveries are automatically calculated by the instrument.

9. REPRESENTATIVE DOCUMENTS

TABLE 3 - Results table / Polar bear liver

	File Name PBISSW3							Cal. Std. File:	3Q quentitation		
			ATVPENPEON		Lipid %		1	PBQ vol. aL		180	1
			5:16 am		Sample Wei		5.23	I.S. (2.5 ng/uL)		10	
	Operator Couriney				New N.F. (1	38)	1.050	E.S. (6 ng/ul.)		10	1
	Method File PBQ_NO		. 1					IS #2 DOE	(2ng/uL)	. 10	1
	Sampie Nam PB X1176 More Inso	ız (m-5,23	»;					SAM F.V. ut.	ON Dr	0. 210	1
	# of Compou 59			* 1	3 , 150			Opini L. V. UC	1.5	210	
	Last Cal. Up: Wed Dec	15 14 11211	9.1000	27 2 2						Ly 1/2	
		STD R.T	STD	STD	SAM R.T.	SAM	conc.	Total	Conc.	Rec.Cor	1
		(min.)	(conc.)	(area)	(m(n.)	(area)	(pg/uL)	Amount	w.w.	1100.00	
		()	(pg/uL)	(e: 02)	(11411.)	(0.00)	(99,00)	(pg)	(pg/g)	(pg/g)	
_	Performance Standard		17-57		-			N-D/	V-07		f
	C13 - PCB138 (ES)	19,95	222.38	205747	19.94	195881					
	Individual CHC Concent		222.00	200741	18.54	100001					1
	TeCIBz	6.69	23	35892	8.73	49653	29	6016	1150.25	1813,19	i
	PnÇIBz	8.48	40	49881	8.53	10435	8	1588	303.72	404,56	
	HC8	10.58	936	938047	10.60	10587	ě	1994	381.17	552.00	
	a-HCH	10.46	114	103218	10.30	19780	20	4130	789.74	1143.70	
,	b-HCH	11.08	29	24732	10.98	20852	22	4623	883.89	1280.03	
	ocs	15.13	14	6614	15,12	2468	5	986	188.85	273.49	
	Compound C	12.23	32	2514	12.24	5438	62	13089	2502.59	3624,23	1
	Photoheplachlor	14.63	49	19211	14.63	9894	22	4675	693.83	1294,44	l
	Heptschlor Epoxide	15.26	199	98191	15 28	4607	16	2367	643,70	\$32,20	1
0	Oxychlorodane	15.30	728	82790	15.29	45815	361	75836	14500,27	20999.13	l
U	⊔-4	15.46	68	14012	15,45	16894	74	15501	2983.87	4292,24	1
2	C-5	10.17	13	10947	16.18	9397	10	2110	403.42	584,23	ı
	C-3	15.84	18	20703	15.84	21663	17	3564	681,51	986,95	1
	C-4	16.46	15	8796	18,45	3321	5	1071	204.73	296.48	
	Nonachior III	16.46	205	132459	18.46	43037	50	12593	2407.83	3486,99	
	- Nonaction	16.64	312	164224	18.64	23421	40	8413	1608,58	2329.49	1
	J-2	18.21	48	13895	18.18	5437	17	3603	688.91	997,67	
	Di-Idrin	17.26	564	20673	17.27	1179	20	4227	810.11	117.3.19	
	p,p'-DDE	17,13	694	860754	17.12	32995	24	5030	961.70	1392.73	
	p,p'-000	18,50	29	33320	18.50	1323	1	218	41.63	60,28	
	p,p'-DDT	19.76	47	67062	19.78	2095	2	384	73.35	106.22	ı
	PCB 52	13.58	n/a	18658	13.54	4073	12	2491	476.28	542,77	
	PCB 47/48	13.77	61	79576	13.77	9881	7	1432	273.82	312.04	
	PCB 74	15.25	37	53400	15.22	4577	3	600	114.64	130.65	
	PCB 56/60	18.02	11	2019	15.99	578	2	426	81.54	92.92	
	PCB 101	16.21	n/a	39028	16.20	8002	10	2132	407.68	464.59	
	PCB 99	16,39	710	746099	16.37	148819	128	26775	5119.56	5834.20	l l
	CB 85	17.23	55	60735	17.21	8249	7	1412	270.04	307.74	}
	CB 149	18.19	13	9552	18.20	3423	4	881	158.41	191,91	I
	PCB 118	1B,26	130	148375	18.25	11463	9	1897	362.76	413.39	
	PCB 146	18.81	59	42472	18.60	8560	11	2248	429.87	489,88	
	PCB 153	19,02	3222	2281430	19.01	518232	659	138375	26457.88	30151,16	
	PC8 105	19.15	36	38852	19.13	4133	4	768 2596	148.75	187.24	
	PCB 137	19.68	43	26211	19.67	8375	12	23180	496,68	566,02	
	PCB 138/163 PCB 167	19.95 20.54	817 40	432934	19,94	85954 3454	110 8	1356	4428,35 259,26	5046,51 295,44	
	CB 183	20.70	38	19265	20.52 20.69		13	2629	502.73	572,91	
	CB 158	21,70	34	190 9 4 27052	21.71	7376 9514	11	2281	432.28	482.62	
	CB 157	21.90	34	25229	21.90	7947	10	2025	387.18	441,21	Ι'
	CB 180	22.28	1225	632830	22.32	160855	280	58797	11242,30	12811.63	l
	CB 170/190	23.28	622	314908	23.44	60614	108	22036	4328.05	4932.21	l
	CB 195	23.78	10	4872	23.90	2211	4	858	164.06	188.06	ı
	CB 194	25.48	212	80695	25.70	10602	25	5352	1023.37	1168,22	
	CB 206	26.77	20	13452	26.95	3388	5	947	181.02	206.29	
	CB 200	27.81	23.5	3587	27.89	769	5	953	182.12	207.55	ł
	encent Recoveries: Inte			9001						rec.com.	1
è			- A						Total	Total	ı
7	ompound INTFILE	TO R.T	STD	STO	SAM R.T.	SAM	Recovery		1835.14	2769.76	sum Cli
	,	(min.)	(conc.)	(area)	(min.)	(area)	%	·	8753	1674	sum HC
			(ng/uL)	,				***	1076.68	1559,23	sum DD
_		***************************************						avgClSz	27044.38	39165.35	sum OC
1	3C-TeClBz	6.69	119.05	107114	6.73	64692	63.4	69.2			1
	3C-PnClBz	8.48	119.05	109594	13.77	78331	75.1		57936.62	86024.04	sum PC
	3C-HCB	10.58	119.05	96841	15.22	63664		•			1
	3C-PCB 28	12.60	93.81	111498	12.59	100293	94.5				I
	3C-PCB 52	13,53	95.23	122901	13.54	103470	88.4	mean PCB			ŀ
	3C-PCB 118	18.26	100.24	118158	18.24	102368	91.0	% recovery			I
	3C-PCB 153	19.02	102.62	79049	19,00	65027	86.4	87.6]
1	3C-PCB 180	22.28	96,1	31567	22.31	28139	93.6				l
	3C-PCB 194	25.48	98.57	39042	25.68	26969	72.6		i		l
	13 - B-RCH	11.05	95.2	24283	10.98	17745		7 12 1	•		l
	13 - DDE	17,16	95.2	135630	17.11	119212	92.3	-			J
C								a(CB154(std.)/C	@ 154(sam.))		l
	18. QUIL, - 1310. QUIL.										
Ī				sample) :	ng						i .
T	otal Amt. = Init. Conc.*(onc. w.w. = (Total Amt.	\$m\$4/5m\$	7)"(F.V. ut c	of sample) =	ng						

10. QUALITY CONTROL

10.1. MSD Tuning

See Section 10.1 of MET-CHEM-OC-04.

10.2. Calibration Verification

A four-point initial calibration standard curve is generated every six months for the major compounds (e.g., oxychlordane, PCB-153, etc.) found in the polar bear quality control material (PBQ) to cover the range of interest. This established calibration curve is verified daily, by analyzing a calibration verification standard (quantitation standard) having a mid-point concentration. The calculated concentration of each compound must be within 20% of its actual known value. The final concentration of any reportable compounds must be within the demonstrated linearity of the detector. If necessary, samples are diluted with iso-octane to meet the calibration range.

10.3. Detection Limit

Minimum detectable concentration (MDC) is the concentration of analyte which produces a signal 3 times the average noise level. Detection limit, using MSD, is highly variable with compound since background noise and response factors are different for each ion monitored. The background noise is affected by several factors, such as tissue size and tissue type, instrument (column bleed and cleanliness of the source on the day of the analysis). Typically a detection limit of 0.001 ppm (or less) is achievable. MDC for individual compounds are calculated periodically using the noise calculation function of the ChemStation software.

10.4. Ongoing Precision and Recovery

An aliquot of a PBQ Reference Material is analyzed along with each batch of samples. The concentration of the major compounds is determined and the results are compared to the previously established acceptance limits (i.e., ± 2 SD of the long-term mean plotted in a Shewart chart - ref. SOP-CHEM-DOC-02).

To determine the degree of analyte loss during sample cleanup, each sample (including the PBQ reference material) is spiked with ¹³C-labeled chlorobenzenes/PCBs internal standard mixture. It is assumed that during

sample cleanup, native compounds behave the same way as the labeled compounds. Analysis is accepted when the % internal standard recoveries for most PCBs and OCs are between 80% and 110%, and for the highly volatile compounds (eluting in the 1st, 2nd and 3rd chromatographic windows) are over 60%. The levels of contaminants reported in the analytical test report are not corrected for internal standard recoveries, but the recovery data are included.

10.5. Accuracy

To improve the accuracy of the method, the compound levels can be corrected based on the recovery of the ¹³C-labeled chlorobenzenes/PCBs internal standard mixture (4.6.1). The majority of the chlorinated pesticides and PCB congeners levels can be corrected based on the average recovery of the five higher chlorinated ¹³C-labeled PCB standards (i.e., PCB-52, PCB-118, PCB-153, PCB-180 and PCB-194). The more volatile compounds can be recovery-corrected using the recovery of their corresponding ¹³C-labeled standards. For volatile compounds without ¹³C-surrogates, retention time interpolation of recoveries with the ¹³C-standards can be used.

The accuracy of the quantitation standards is verified annually with a second source standard (containing most of the congeners of interest) as described in SOP-CHEM-PROC-13.

10.6. Method Blank

See Section 10.6 of MET-CHEM-OC-04.

10.7. Target Compound Identification

See Section 10.7 of MET-CHEM-OC-04.

10.8. Standard Operating Procedures

See Section 10.8 of MET-CHEM-OC-04.

10.9. Data Validation

See Section 10.9 of MET-CHEM-OC-04.

10.10. External Quality Control

As part of its quality-assurance effort, the Laboratory organized an interlaboratory study involving the determination of 33 compounds in a polar bear fat sample. Results obtained in our laboratory were in good agreement with the results in the other five participant laboratories (average of 35% variation for the 33 compounds examined) [2.9].

11. CRITICAL CONTROL POINTS

See Section 11 of MET-CHEM-OC-04 and:

The flow of nitrogen (used to concentrate the extracts) has to be adjusted so that the surface of the solvent is just visibly disturbed and the sample must never be brought to dryness as it may cause analyte loss.

Analysts must be aware that electronic integration of data has its pitfalls. Examination of chromatograms and results by a competent analyst is essential.

When doing timed activities ensure each sample is treated equally (e.g., when doing moisture or lipid determinations).

APPENDIX A

Numbering of PCB Congeners

Numbering of PCB congeners - Adapted from Ballschmiter, K. et al. (1980) Anal. Chem., 302, 20-31.

	Monochlorobiphenyls		<u>Tetrachlorobiphenyls</u>		Pentachlorobiphenyls	Hexachlorobiohenyls	
1	2	52	2,2',5,5'	105	2,3,3',4,4'	161	2,3,3',4,5',6
2	3	53	2,2',5,6	106	2,3,3',4,5	162	2,3,3',4',5,5'
3		54	2,2',6,6'	107		163	2,3,3',4',5,6
3	4				2,3,3',4',5	164	
	Bill bishawk	55	2,3,3',4	108	2,3,3',4,5'		2,3,3',4',5',6
	<u>Dichlorobiphenyls</u>	56	2,3,3',4'	109	2,3,3',4,6	165	2,3,3',5,5',6
		57	2,3,3',5	110	2,3,3',4',6	166	2,3,4,4',5,6
4	2,2'	58	2,3,3',5'	111	2,3,3',5,5'	167	2,3',4,4',5,5'
5	2,3	59	2,3,3',6	112	2,3,3',5,6	168	2,3',4,4',5',6
6	2,3'	60	2,3,4,4'	113	2,3,3',5',6	169	3,3',4,4',5,5'
7	2,4	61	2,3,4,5	114	2,3,4,4',5		
8	2,4'	62	2,3,4,6	115	2,3,4,4',6		<u>Heptachlorobiphenyls</u>
9	2,5	63	2,3,4',5	116	2,3,4,5,6		
10	2,6	64	2,3,4',6	117	2,3,4',5,6	170	2,2',3,3',4,4',5
11	3,3'	65	2,3,5,6	118	2,3',4,4',5	171	2,2',3,3',4,4',6
12	3,4	66	2,3',4,4'	119	2,3',4,4',6	172	2,2',3,3',4,5,5'
13	3.4'	67	2,3',4,5	120	2,3',4,5,5'	173	2,2',3,3',4,5,6
14	3,5	68	2,3',4,5'	121	2,3',4,5',6	174	2,2',3,3',4,5,6'
		69		122		175	
15	4, 4'		2,3',4,6		2',3,3',4,5		2,2',3,3',4,5',6
		70	2,3',4',5	123	2',3,4,4',5	176	2,2',3,3',4,6,6'
	<u>Trichlorobiphenvis</u>	71	2,3',4',6	124	2',3,4,5,5'	177	2,2',3,3',4',5,6
		72	2,3',5,5'	125	2',3,4,5,6'	178	2,2',3,3',5,5',6
16	2,2',3	73	2,3',5',6	126	3,3',4,4',5	179	2,2',3,3',5,6,6'
17	2,2',4	74	2,4,4,5	127	3,3',4,5,5'	180	2,2',3,4,4',5,5'
18	2,2',5	75	2,4,4',6		,	181	2,2',3,4,4',5,6
19	2,2',6	76	2,3,4,5		Hexachlorobiphenyls	182	2,2',3,4,4',5,6'
20	2,3,3'	77	3,3',4,4'		T TO A CONTROL OF THE PARTY OF	183	2,2',3,4,4',5',6
		78	3,3',4,5	128	2 21 2 21 4 41	184	2,2',3,4,4',6,6'
21	2,3,4	79		129	2,2',3,3',4,4'	185	
22	2,3,4		3,3',4,5'		2,2',3,3',4,5		2,2',3,4,5,5',6
23	2,3,5	80	3,3',5,5'	130	2,2',3,3',4,5'	186	2,2',3,4,5,6,6'
24	2,3,6	81	3,4,4',5	131	2,2',3,3',4,6	187	2,2',3,4',5,5',6
25	2,3',4			132	2,2',3,3',4,6'	188	2,2',3,4',5,6,6'
26	2,3',5		Pentachlorobiphenyls	133	2,2',3,3',5,5'	189	2,3,3',4,4',5,5'
27	2,3',6			134	2,2',3,3',5,6	190	2,3,3',4,4',5,6
28	2,4,4'	82	2,2',3,3',4	135	2,2',3,3',5,6'	191	2,3,3',4,4',5',6
29	2.4.5	83	2,2',3,3',5	136	2,2',3,3',6,6'	192	2,3,3',4,5,5',6
30	2.4.6	84	2,2',3,3',6	137	2,2',3,4,4',5	193	2,3,3',4',5,5',6
31	2,4',5	85	2,2',3,4,4'	138	2,2',3,4,4',5'		_, , , . , . , . , . , . , . , . , . , .
32	2.4'.6	86	2,2',3,4,5	139	2,2',3,4,4',6		Octachlorobiphenyls
33	2',3,4	87	2,2',3,4,5'	140	2,2',3,4,4',6'		Octabilioropiphertyrs
33 34		88		141		194	2 21 2 21 4 41 6 61
	2',3,5		2,2',3,4,6		2,2',3,4,5,5'		2,2',3,3',4,4',5,5'
35	3,3',4	89	2,2',3,4,6'	142	2,2,3,4,5,6	195	2,2',3,3',4,4',5,6
36	3,3',5	90	2,2',3,4',5	143	2,2',3,4,5,6'	196	2,2',3,3',4,4',5,6'
37	3,4,4'	91	2,2',3,4',6	144	2,2',3,4,5',6	197	2,2',3,3',4,4',6,6'
38	3,4,5	92	2,2',3,5,5'	145	2,2',3,4,6,6'	198	2,2',3,3',4,5,5',6
39	3,4,5	93	2,2,3,5,6	146	2,2',3,4',5,5'	199	2,2',3,3',4,5,6,6'
		94	2,2',3,5,6'	147	2,2',3,4',5,6	200	2,2',3,3',4,5',6,6'
	<u>Tetrachlorobiphenyls</u>	95	2,2',3,5',6	148	2.2',3,4',5,6'	201	2,2',3,3',4,5,5',6'
		96	2,2',3,6,6'	149	2,2',3,4',5',6	202	2,2',3,3',5,5',6,6'
40	2,2',3,3'	97	2,2',3',4,5	150	2,2',3,4',6,6'	203	2,2',3,4,4',5,5',6
41	2,2',3,4	98	2,2',3',4,6	151	2,2',3,5,5',6	204	2,2',3,4,4',5,6,6'
42	2.2'.3.4'	99	2,2',4,4',5	152	2,2',3,5,6,6'	205	2,3,3',4,4',5,5',6
						200	2,3,3,4,4,3,3,6
43	2,2,3,5	100	2,2',4,4',6	153	2,2',4,4',5,5'		Managhtagettetered
44	2,2',3,5'	101	2,2',4,5,5'	154	2,2',4,4',5,6'		Nonachlorobiohenyls
45	2,2',3,6	102	2,2',4,5,6'	155	2,2',4,4',6,6'		
46	2,2',3,6'	103	2,2',4,5',6	156	2,3,3',4,4',5	206	2,2',3,3',4,4',5,5',6
47	2,2',4,4'	104	2,2',4,6,6'	157	2,3,3',4,4',5'	207	2,2',3,3',4,4',5,6,6'
48	2,2',4,5			158	2,3,3',4,4',6	208	2,2,3,3,4,5,5,6,6
49	2,2,4,5			159	2,3,3',4,5,5'		
50	2,2',4,6			160	2,3,3',4,5,6		<u>Decachlorobiphenyl</u>
51	2,2',4,6'						
٥.	-1-1 1 V					209	2,2',3,3',4,4',5,5',6,6'
						200	51210121210101010