

OptiPrep™ Mini-Review MS02

Purification of peroxisomes – a bibliography

There are several OptiPrep™ Application Sheets that are relevant to the isolation and analysis of peroxisomes using iodixanol gradients and three types of gradient have been used:

Continuous iodixanol gradients: principally for the purification of peroxisomes from mammalian liver, but they have also been used for organelles from fungi, mainly yeast and also from plants.

Discontinuous iodixanol gradients: for mammalian liver and kidney, some marine organisms and plants.

Self-generated iodixanol gradients: for mammalian liver and cultured cells

- ◆ **Application Sheet S11** describes the use of a pre-formed continuous iodixanol gradient
- ◆ **Application Sheet S12** describes the use of a discontinuous iodixanol gradient
- ◆ **Application Sheet S13** describes the use of a self-generated iodixanol gradient
- ◆ **Application Sheet S55** describes the use of a continuous iodixanol gradient for yeast peroxisomes

In addition there are two Application Sheets that are devoted to the analysis of the light mitochondrial fraction (LMF) and although these are not devoted specifically to peroxisomes, the latter are analyzed as part of a more general analysis of the LMF organelles, which include mitochondria, lysosomes and sometimes Golgi in addition to peroxisomes.

- ◆ **Application Sheet S15** describes the use of pre-formed continuous gradient
- ◆ **Application Sheet S16** describes the use of self-generated gradient

The bibliography below is divided into gradient type, then tissue or cell source. References are listed alphabetically according to **first author** and then, if required, chronologically. To aid identification of research topics, these are highlighted in blue.

1. Continuous gradients

1a. Brain (rodent)

Nawrotzki, R., Islinger, M., Vogel, I., Völkl, A. and Kirsch, J. (2012) *Expression and subcellular distribution of gephyrin in non-neuronal tissues and cells* Histochem. Cell. Biol., **137**, 471–482

1b. Fat pad (mammary)

Vapola, M.H., Rokka, A., Sormunen, R.T., Alhonen, L., Schmitz W., Conzelmann, E., Wärri, A., Grunau, S., Antonenkov, V.D. and Hiltunen, J.K. (2014) *Peroxisomal membrane channel Pxmp2 in the mammary fat pad is essential for stromal lipid homeostasis and for development of mammary gland epithelium in mice* Dev. Biol., **391**, 66–80

1c. Fibroblasts

Wiesinger, C., Kunze, M., Regelsberger, G., Forss-Petter, S. and Berger, J. (2013) *Impaired very long-chain Acyl-CoA β-oxidation in human X-linked adrenoleukodystrophy fibroblasts is a direct consequence of ABCD1 transporter dysfunction* J. Biol. Chem., **288**, 19269–19279

1d. Fungi

1d-1 *Paracoccidioides brasiliensis*

Brito, W.deA., Rezende, T.C.V., Parente, A.F., Ricart, C.A.O., de Sousa, M.V., Bão, N. and Soares, C.M.deA. (2011) *Identification, characterization and regulation studies of the aconitase of Paracoccidioides brasiliensis* Fungal Biol., **115**, 697–707

1d-2 Yeast

- Antonenkov, V.D.**, Mindthoff, S., Grunau, S., Erdmann, R. and Hiltunen, J.K. (2009) *An involvement of yeast peroxisomal channels in transmembrane transfer of glyoxylate cycle intermediates* Int. J. Biochem. Cell Biol., **41**, 2546–2554
- Cramer, J.**, Effelsberg, D., Girzalsky, W. and Erdmann, R. (2015) *Isolation of peroxisomes from yeast* Cold Spring Harb. Protoc; doi:10.1101/pdb.top074500
- Cramer, J.**, Effelsberg, D., Girzalsky, W. and Erdmann, R. (2015) *Small-scale purification of peroxisomes for analytical applications* Cold Spring Harb. Protoc; doi:10.1101/pdb.prot083717
- Debelyy, M.O.**, Platta, H.W., Saffian, D., Hensel, A., Thoms, S., Meyer, H.E., Warscheid, B., Girzalsky, W. and Erdmann, R. (2011) *Ubp15p, a ubiquitin hydrolase associated with the peroxisomal export machinery* J. Biol. Chem., **286**, 28223–28234
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- Grunau, S.**, Mindthoff, S., Rottensteiner, H., Sormunen, R.T., Hiltunen, J.K., Erdmann, R. and Antonenkov, V.D. (2009) *Channel-forming activities of peroxisomal membrane proteins from the yeast Saccharomyces cerevisiae* FEBS J., **276**, 1698–1708
- Grunau, S.**, Lay, D., Mindthoff, S., Platta, H.W., Girzalsky, W., Just, W.W. and Erdmann, R. (2011) *The phosphoinositide 3-kinase Vps34p is required for pexophagy in Saccharomyces cerevisiae* Biochem. J. **434**, 161–170
- Kerssen, D.**, Hambruch, E., Klaas, W., Platta, H.W., de Kruijff, B., Erdmann, R., Kunau, W-H. and Schliebs, W. *Membrane association of the cycling peroxisome import receptor Pex5p* J. Biol. Chem., **281**, 27003-27015
- Mindthoff, S.**, Grunau, S., Steinfort, L.L., Girzalsky, W., Hiltunen, J.K., Erdmann, R. and Antonenkov, V.D. (2016) *Peroxisomal Pex11 is a pore-forming protein homologous to TRPM channels* Biochim. Biophys. Acta, **1863**, 271–283
- Oeljeklaus, S.**, Reinartz, B.S., Wolf, J., Wiese, S., Tonillo, J., Podwojski, K., Kuhlmann, K., Stephan, C., Meyer, H.E., Schliebs, W., Brocard, C., Erdmann, R. and Warscheid, B. (2012) *Identification of core components and transient interactors of the peroxisomal importomer by dual-track stable isotope labeling with amino acids in cell culture analysis* J. Proteome Res. 2012, 11, 2567–2580
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1e. Liver (rodent)

- Antonenkov, V.D.**, Sormunen, R.T. and Hiltunen, J.K. (2004) *The behavior of peroxisomes in vitro: mammalian peroxisomes are osmotically sensitive particles* Am. J. Physiol., **287**, C1623-C16350
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- Antonenkov, V.D.**, Sormunen, R.T., Ohlmeier, S., Amery, L., Fransen, M., Mannaerts, G.P. and Hiltunen, J.K. (2006) *Localization of a portion of the liver isoform of fatty-acid-binding protein (L-FABP) to peroxisomes* Biochem. J., **394**, 475-484
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- Graham, J.**, Ford, T. and Rickwood, D (1994) *The preparation of subcellular organelles from mouse liver in self-generated gradients of iodixanol* Anal. Biochem., **220**, 367-373

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1f. Mammary fat pad

Vapola, M.H., Rokka, A., Sormunen, R.T., Alhonen, L., Schmitz W., Conzelmann, E., Wärri, A., Grunau, S., Antonenkov, V.D. and Hiltunen, J.K. (2014) *Peroxisomal membrane channel Pxmp2 in the mammary fat pad is essential for stromal lipid homeostasis and for development of mammary gland epithelium in mice* Dev. Biol., **391**, 66–80

1g. Plant tissues

- Arai, Y.**, Hayashi, M. and Nishimura, M. (2008) *Proteomic analysis of highly purified peroxisomes from etiolated Soybean cotyledons* Plant Cell Physiol., **49**, 526-539
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1h. Review

Antonenkov, V.D. and Hiltunen, J.K. (2006) *Peroxisomal membrane permeability and solute transfer* Biochim. Biophys. Acta, Mol. Cell Res., **1763**, 1697-1706

2. Discontinuous gradients

2a. HEK cells

Ge, L., Melville, D., Zhang, M. and Schekman, R. (2013) *The ER–Golgi intermediate compartment is a key membrane source for the LC3 lipidation step of autophagosome biogenesis* eLife, **2**: e00947

Zhang, J., Kim, J., Alexander, A., Cai, S., Tripathi, D.N., Dere, R., Tee, A.R., Tait-Mulder, J., Di Nardo, A., Han, J.M., Kwiatkowski, E., Dunlop, E.A., Dodd, K.M., Folkerth, R.D., Faust, P.L., Kastan, M.B., Sahin, M. and Walker, C.L. (2013) *A tuberous sclerosis complex signalling node at the peroxisome regulates mTORC1 and autophagy in response to ROS* Nat. Cell Biol., **15**, 1186-1196

2b. Human fibroblasts

Beltran, P.M.J., Mathias, R.A. and Cristea, I.M. (2016) *A portrait of the human organelle proteome in space and time during cytomegalovirus infection* Cell Systems **3**, 361–373

2c. Kidney (rodent)

Mi, J., Garcia-Arcos, I., Alvarez, R., and Cristobal, S. (2007) *Age-related subproteomic analysis of mouse liver and kidney peroxisomes* Proteome Sci., **5:19**

Mi, J., Kirchner, E. and Cristobal, S. (2007) *Quantitative proteomic comparison of mouse peroxisomes from liver and kidney* Proteomics, **7**, 1916-1928

2d. Liver (chick embryo)

Labitzke, E.M., Diani-Moore, S. and Rifkind, A.B. (2007) *Mitochondrial P450-dependent arachidonic acid metabolism by TCDD-induced hepatic CYP1A5; conversion of EETs to DHETs by mitochondrial soluble epoxide hydrolase* Arch. Biochem. Biophys., **468**, 70-81

2e. Liver (rodent)

Amelina, H., Sjödin, M.O.D., Bergquist, J. and Cristobal, S. (2011) *Quantitative subproteomic analysis of age-related changes in mouse liver peroxisomes by iTRAQ LC-MS/MS* J. Chromatogr. B, **879**, 3393– 3400

Grant, P., Ahlemeyer, B., Karnati, S., Berg, T., Stelzig, I., Nenicu, A., Kuchelmeister, K., Crane, D.I. and Baumgart-Vogt, E. (2013) *The biogenesis protein PEX14 is an optimal marker for the identification and localization of peroxisomes in different cell types, tissues, and species in morphological studies* Histochem. Cell. Biol., **140**, 423–442

Karnati, S., Lüers, G., Pfreimer, S. and Baumgart-Vogt, E. (2013) *Mammalian SOD2 is exclusively located in mitochondria and not present in peroxisomes* Histochem. Cell Biol., **140**, 105–117

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Salvi, M., Battaglia, V., Brunati, A.M., La Rocca, N., Tibaldi, E., Pietrangeli, P., Marcoucci, L., Mondovi, B., Rossi, C.A. and Toninello, A. (2007) *Catalase takes part in rat liver mitochondria oxidative stress defense* J. Biol. Chem., **282**, 24407-24415

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2f. Mouse embryo fibroblasts

Zhang, J., Kim, J., Alexander, A., Cai, S., Tripathi, D.N., Dere, R., Tee, A.R., Tait-Mulder, J., Di Nardo, A., Han, J.M., Kwiatkowski, E., Dunlop, E.A., Dodd, K.M., Folkerth, R.D., Faust, P.L., Kastan, M.B., Sahin, M. and Walker, C.L. (2013) *A tuberous sclerosis complex signalling node at the peroxisome regulates mTORC1 and autophagy in response to ROS* Nat. Cell Biol., **15**, 1186-1196

2g. Mussels

- Apraiz, I.**, Mi, J. and Cristobal, S. (2006) *Identification of proteomic signatures of exposure to marine pollutants in mussels (*Mytilus edulis*)* Mol. Cell. Proteom., **5**, 1274-1285
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2h. Yeast

- Nyathi, Y.**, De Marcos Lousa, C., van Roermund, C.W., Wanders, R.J.A., Johnson, B., Baldwin, S.A., Theodoulou, F.L. and Baker, A. (2010) *The *Arabidopsis* peroxisomal ABC transporter, Comatose, complements the *Saccharomyces cerevisiae pxa1 pxa2Δ* mutant for metabolism of long-chain fatty acids and exhibits fatty acyl-CoA-stimulated ATPase activity* J. Biol., Chem., **285**, 29892–29902
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3. Self-generated gradient

3a. CHO cells

- Honsho, M.**, Yagita, Y., Kinoshita, N. and Fujiki, Y. (2008) *Isolation and characterization of mutant animal cell line defective in alkyl-dihydroxyacetonephosphate synthase: Localization and transport of plasmalogens to post-Golgi compartments* Biochim. Biophys. Acta, **1783**, 1857-1865
- Kobayashi, S.**, Tanaka, A. and Fujiki, Y. (2007) *Fis1, DLP1 and Pex11p coordinately regulate peroxisome morphogenesis* Exp. Cell Res., **313**, 1675-1686
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3b. Hep-G2 cells

- Morel, F.**, Rauch, C., Petit, E., Piton, A., Theret, N., Coles, B. and Guillouzo, A. (2004) *Gene and protein characterization of the human glutathione S-transferase kappa and evidence for a peroxisomal localization* J. Biol. Chem., **279**, 16246-16253

3c. Liver (rodent)

- He, D.**, Barnes, S. and Falany, C.N. (2003) *Rat liver bile acid CoA:amino acid N-acyltransferase: expression, characterization, and peroxisomal localization* J. Lipid Res., **44**, 2242-2249
- Graham, J.**, Ford, T. and Rickwood, D (1994) *The preparation of subcellular organelles from mouse liver in self-generated gradients of iodixanol* Anal. Biochem., **220**, 367-373
- Kurochkin, I.V.**, Mizuno, Y., Konagaya, A., Sakaki, Y., Schonbach, C. and Okazaki, Y. (2007) *Novel peroxisomal protease Tysnd1 processes PTS1- and PTS2-containing enzymes involved in β-oxidation of fatty acids* EMBO J., **26**, 835-845
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- Miyata, N.**, Hosoi, K-i., Mukai, S. and Fujiki, Y. (2009) *In vitro import of peroxisome-targeting signal type 2 (PTS2) receptor Pex7p into peroxisomes* Biochim. Biophys. Acta, **1793**, 860–870
- Styles, N.A.**, Falany, J.L., Barnes, S. and Falany, C.N. (2007) *Quantification and regulation of the subcellular distribution of bile acid coenzyme A:amino acid N-acyltransferase activity in rat liver* J. Lipid Res., **48**, 1305-1315

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