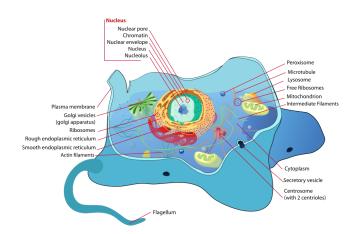
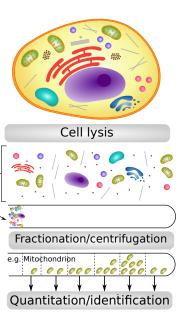
Computational Challenges in Mass Spectrometry-Based Spatial Proteomics

Laurent Gatto
University of Cambridge
http://cpu.sysbiol.cam.ac.uk

8 October 2014





	Fraction ₁	Fraction ₂		$Fraction_m$		markers	
$prot_1$	$q_{1,1}$	$q_{1,2}$		q _{1, m}		unknown	
$prot_2$	$q_{2,1}$	$q_{2,2}$		$q_{2, m}$		$organelle_1$	
$prot_3$	$q_{3,1}$	$q_{3,2}$		$q_{3, m}$		unknown	
prot_4	$q_{4,1}$	$q_{4,2}$		$q_{4, m}$		$organelle_2$	
:	:	:	:	:	:	:	:
$\mathrm{prot}_{\mathrm{i}}$	$q_{i,1}$	$q_{i,2}$		$q_{i, m}$		$organelle_k$	
:	:	:	:	:	:	:	:
$\mathrm{prot}_{\mathrm{n}}$	$q_{n,1}$	$q_{n,2}$		$q_{n, m}$		unknown	
	Fraction ₁	Fraction ₂		$Fraction_m$			
	:	:	:	:			

Figure: Spatial proteomics data and metadata.

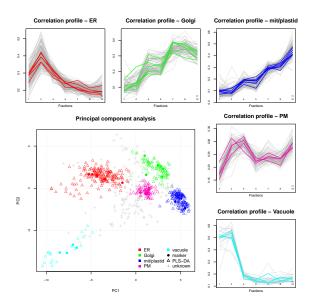


Figure: Visualisation of protein profiles, Gatto et al. (2010).

Challenge: resolution vs. missing data

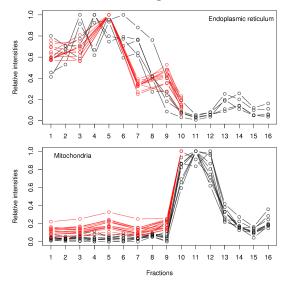


Figure: More fractions of more proteins? Gatto et al. (2014).

Challenge: resolution vs. missing data

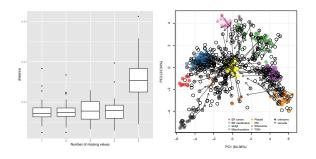


Figure: Missing value imputation. Gatto et al. (2014).

Challenge: spatial markers

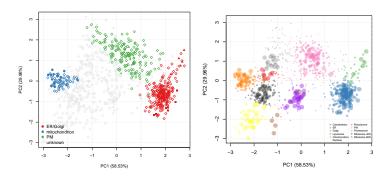
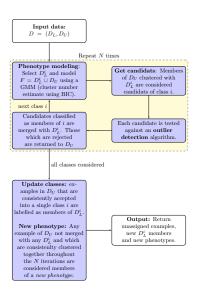


Figure: Sub-cellular diversity. Tan et al. (2009) vs. Breckels et al. (2013).



Challenge: spatial markers

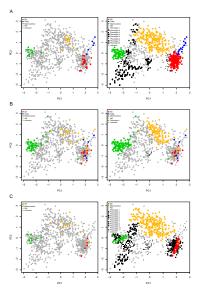


Figure: Quality of markers. Gatto et al. (2014).

Challenge: dual localisation

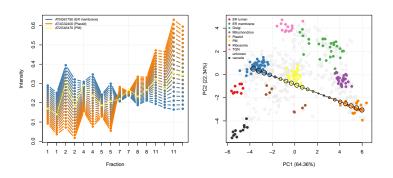


Figure: Gatto et al. (2014).

Challenge: dynamics

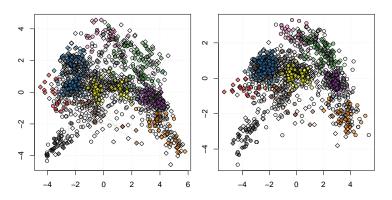


Figure: Normalisation. Gatto et al. (2014).

Challenge: dynamics

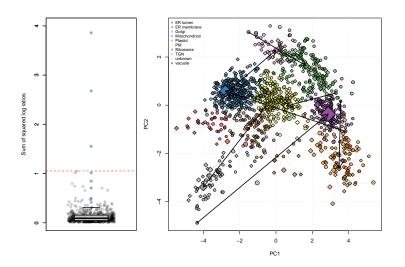
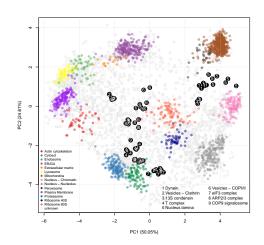
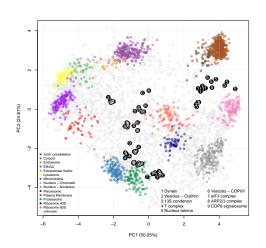


Figure: Statistics. Gatto et al. (2014).

Conclusions



Conclusions



- ► Mass spectrometry
- Experimental design
- ► Computational tools

- ► Funding: BBSRC and FP7 Prime-XS
- Kathryn Lilley
- Lisa Breckels



Gatto L et al. Organelle proteomics experimental designs and analysis. Proteomics 2010 10(22):3957-69. PMID: 21080489.



Gatto L et al. The effect of organelle discovery upon sub-cellular protein localisation. J Proteomics 2013 88:129-40. PMMID 23523639.



Gatto L et al. A foundation for reliable spatial proteomics data analysis. Mol Cell Proteomics 2014 13(8):1937-52. PMID: 24846987.



Gatto L et al. Mass-spectrometry-based spatial proteomics data analysis using pRoloc and pRolocdata. Bioinformatics 2014 30(9):1322-4. PMID: 24413670.



Gatto L et al. A unifying bioinformatics framework for spatial proteomics. Bioconductor 2012 http://www.bioconductor.org/package/release/bioc/html/pRoloc.html