

What the resource is, how it is used, & opportunities for future use

The CCV Prostate Cancer Resource

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Resources

- Melbourne Collaborative Cohort Study: 1990-1994
 - Cohort study Melbourne
 - 17,045 men recruited
- RFPCS: 1994 1998
 - Case-control study Melbourne, Perth & Sydney
 - Final case criteria: Aged ≤70 years + Gleason score >4
- EOPCFS: 1998 (1996) 2008
 - Family-based study Victoria
 - Aged ≤70 years, recruited affected relatives where possible
- APC: 2010 ongoing
 - Case-control study Melbourne
 - Final case criteria: Aged < 75 years + GS >7/TNM T3+/PSA >19
 - Final control criteria: Aged <77 years + biopsy-screened negative

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Resources

		MCCS	RFPCS	EOPCFS	APC
Peripheral Blood	Cases	1,757	1,360	2,269	1,129
	Controls	Matched	1,507		938
Tumour Tissue	Diagnostic	405	247	156	0
	RP	162	133	168	0
	Other	42	42	12	0

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- Genomic determinants of risk and outcomes
 - Peripheral blood
 - Fine-mapping
 - Candidate genes
 - OncoArray
 - Methylation
 - Whole-exome sequencing
 - Tissue
 - IHC Biomarkers PCSM cases + referents
 - Gene expression PCSM cases + referents
 - Methylation PCSM cases + referents

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Collaborations

PRACTICAL

- iSELECT: 47,120 SNPs in 1,818 ca + 1,540 co (M/R/E)
- OPA: 178 SNPs in 1,618 ca + 1,533 co (M/R/E)
- iCOGS: 1,813 ca + 1,231 co (M/R/E)
- OncoArray: 772 ca + 331 co (M)

ICPCG

- Linkage: ?
- Whole-exome sequencing: 15 cases from 11 families
- Omni5Exome: 169 familial ca + 168 RFPCFS co
- Movember (Fred Hutchinson)
 - 30 PCSM-associated SNPs: MCCS and EOPCFS
 - EOPCFS: ~1,670 cases 100 PCSM cases
 - MCCS: ~1,500 cases 199 PCSM cases

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Opportunities for future use

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- Expand whole-exome/genome sequencing
 - More families
 - Tumour tissue
- Expand OncoArray genotyping
 - More than MCCS
- Expand methylation/gene expression efforts
 - More than MCCS (peripheral samples)
 - More than PCSM cases/referents (tumour tissue)
- Tumour tissue collection
 - Further opportunities for collaborative efforts?
 - TMAs?
 - TMPRSS2-ERG?
- Other opportunities?