

Genomics, Proteomics, Transcriptomics

– a Systems Biology Approach to CKD

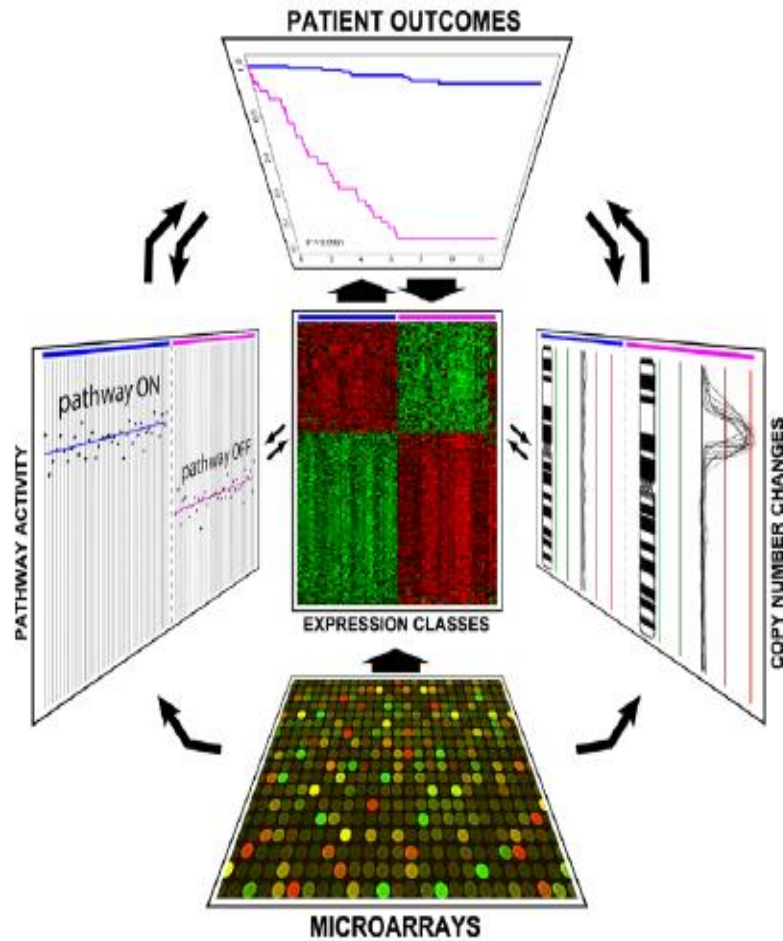
Rainer Oberbauer (30+10min)

What is ,Omics‘?

- Neologism ‘omics’ informally refers to a field of study ending in -omics, such as genomics or proteomics.

- All constituents considered collectively

What is Systems Biology?



Genomics



Genomics





Get to know you. Health and ancestry start here.



- Find out what your DNA says about you
- Get reports on 240+ health conditions & traits
- Read recommendations on steps you can take

[order now](#) **\$99**

Watch a life-changing story.



What your DNA says about you.

Find out things like if your body metabolizes caffeine quickly, or if you're at a higher risk for diabetes. The more you know about your DNA, the more you know about yourself.



Carrier status

Find out if your children are at risk for inherited conditions, so you can plan for the health of your family.



Health risks

Understand your genetic health risks. Change what you can, manage what you can't.



Drug response

Arm your doctor with information on how you might respond to certain medications.

order now



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Mission Statement

23andMe's mission is to be the world's trusted source of personal genetic information.

A Snapshot of Team 23andMe



Who We Are

Board of Directors

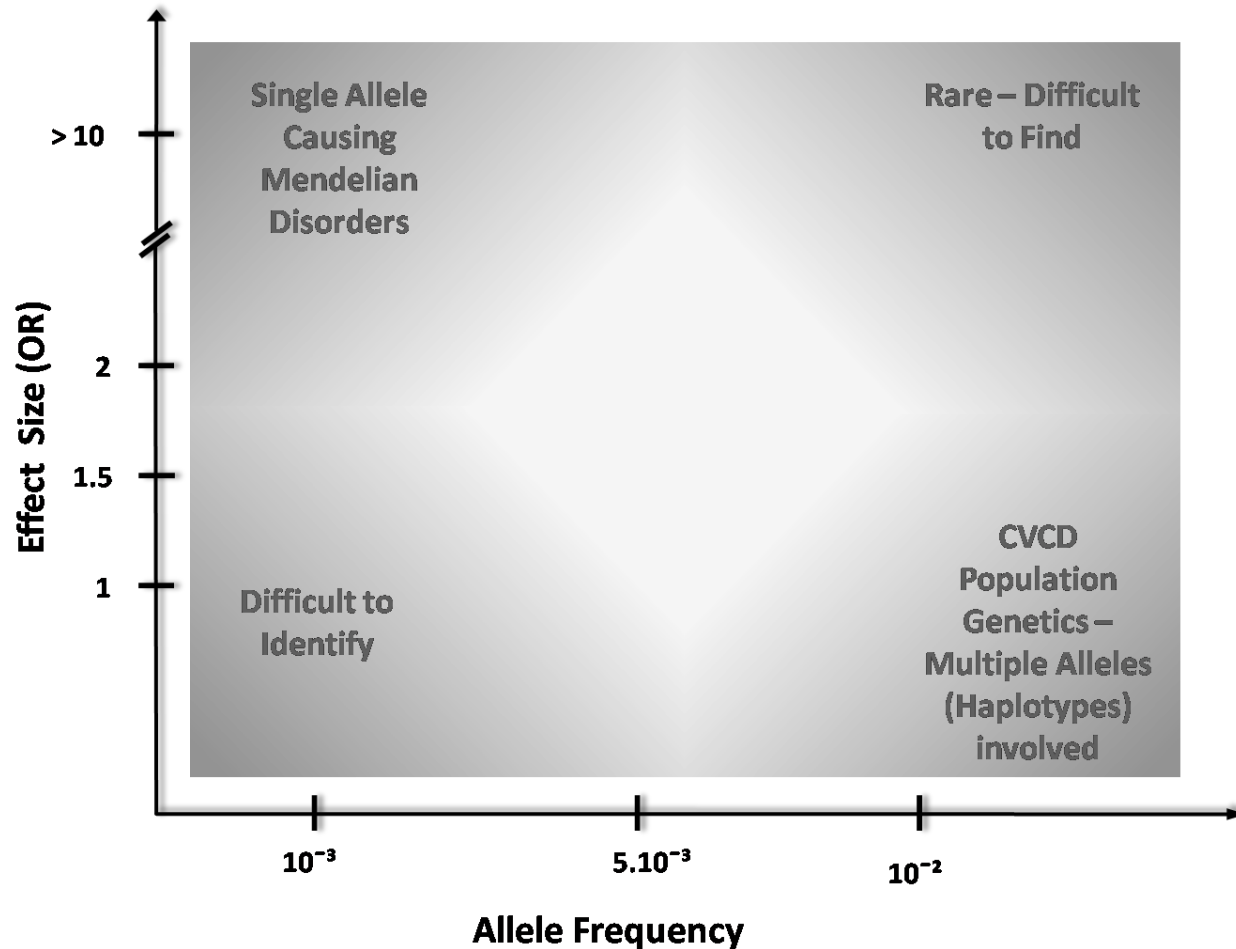
Editorial Team

Advisors

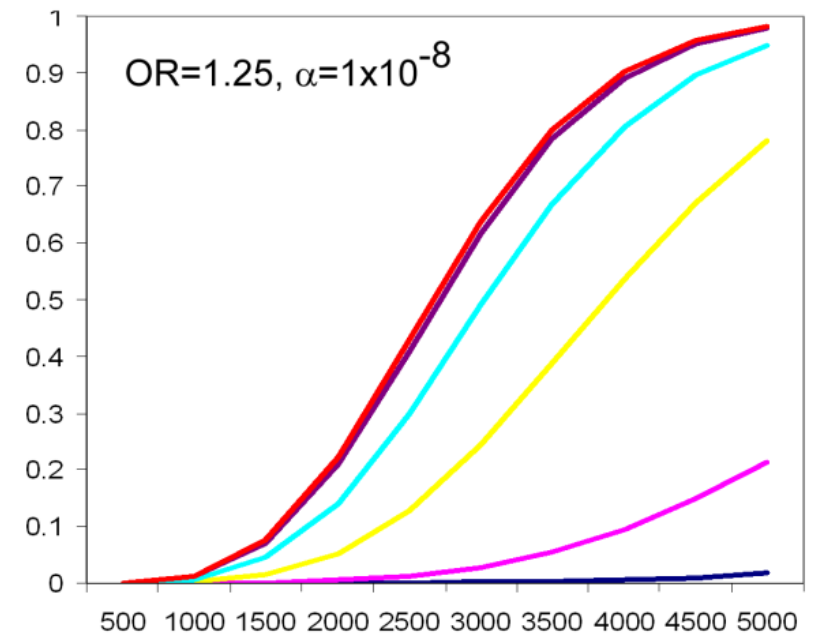
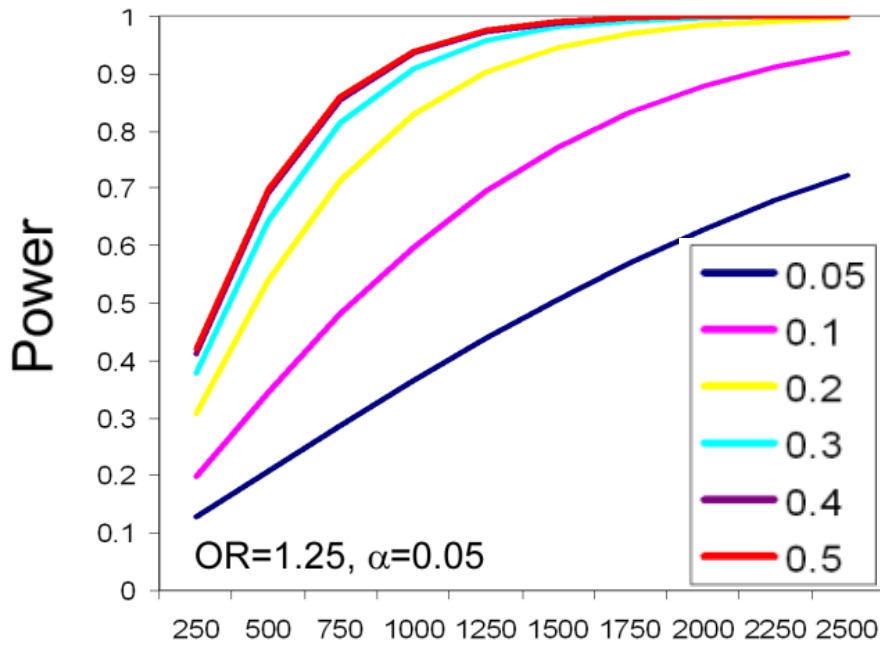
Team 23andMe

67% of us have a family history of cancer

Allele frequencies and effect sizes of genetic variants in human diseases - GWAS

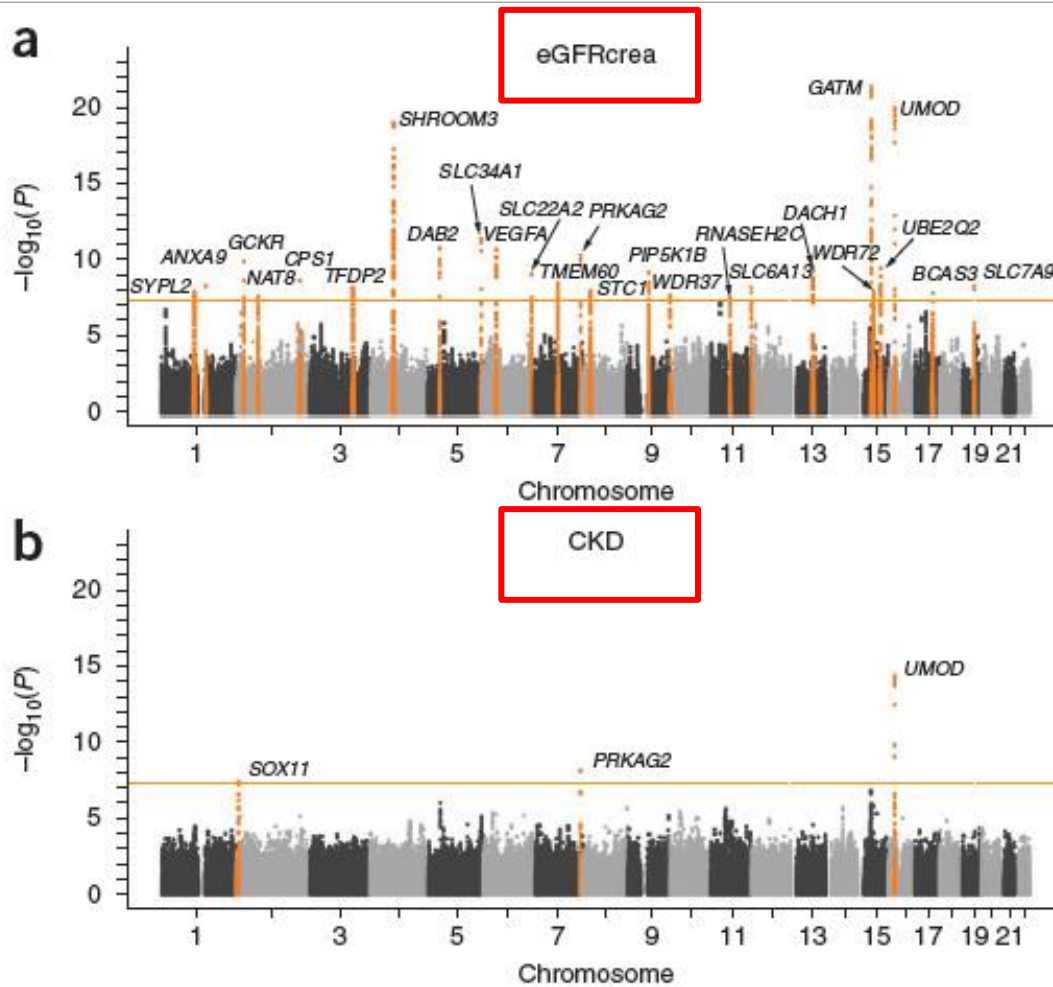


Sample size/power in GWAS



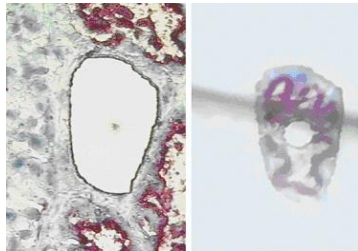
Sample Size

GWAS and CKD

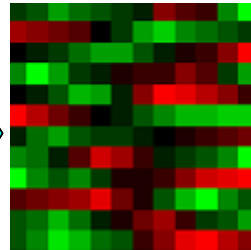


Systems biology of kidney injury

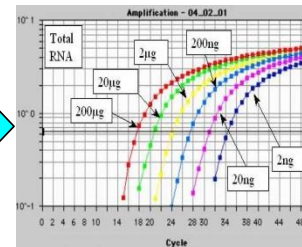
Laser Capture
Microdissection
Of Donor Kidneys



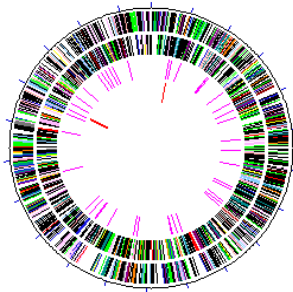
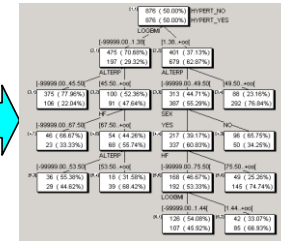
DNA
Microarrays



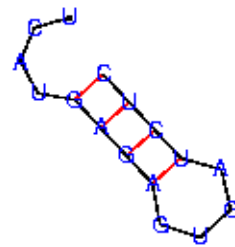
Quantitative
PCR



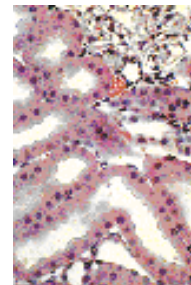
Data
Mining



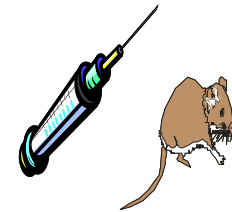
Target Gen
Definition



Antisense
Nucleotide
Design &
Synthese

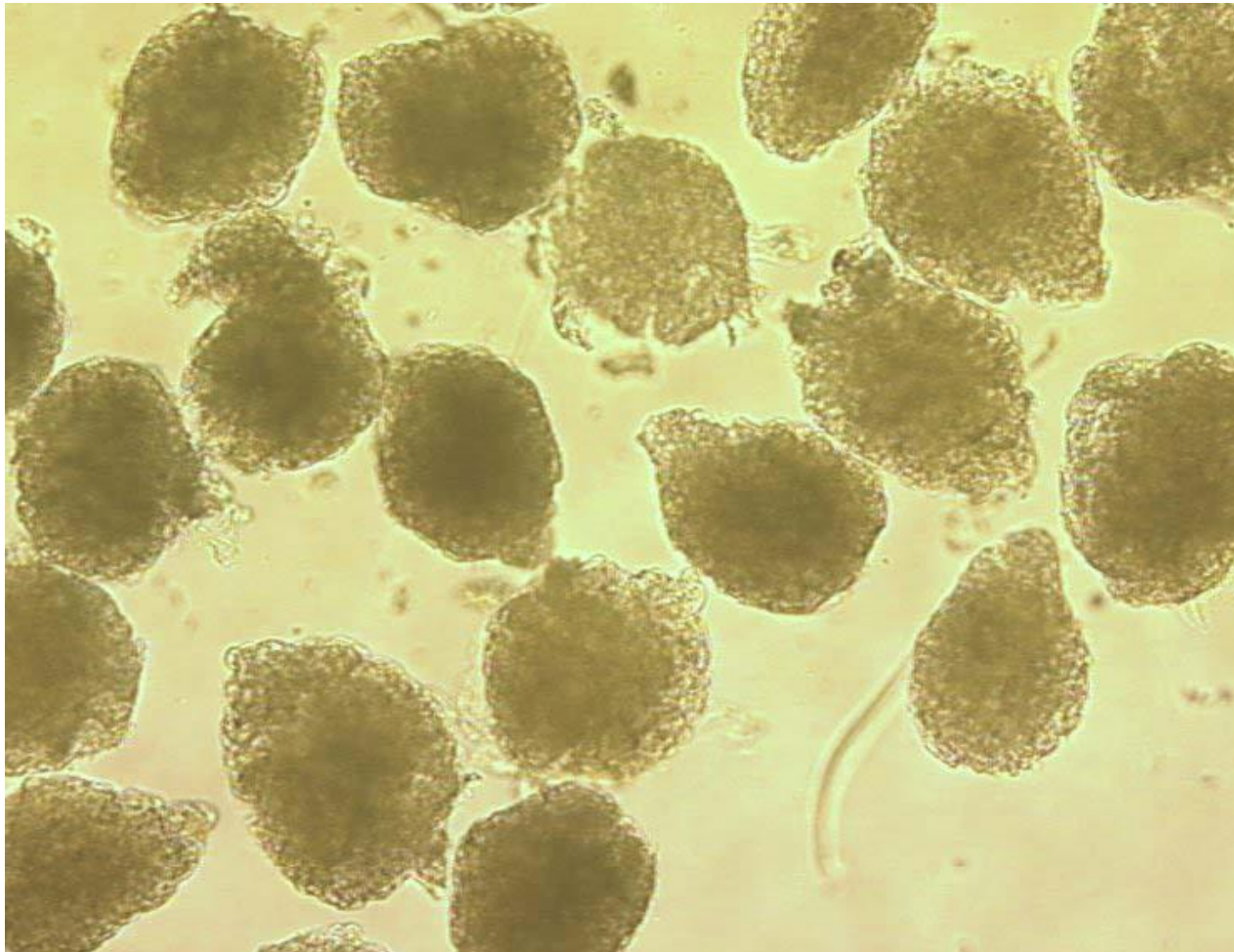


In vitro
POC

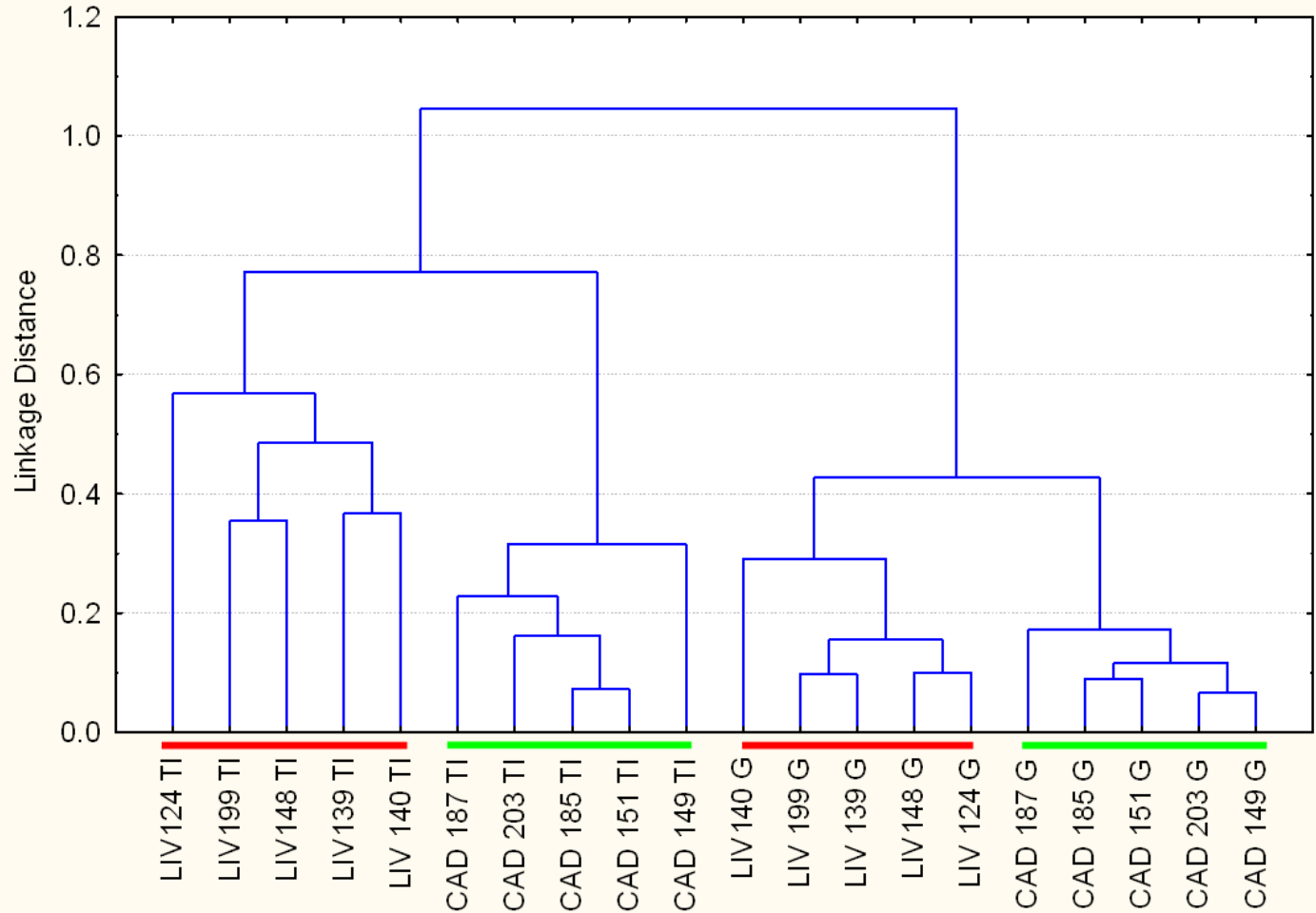


In vivo
POC

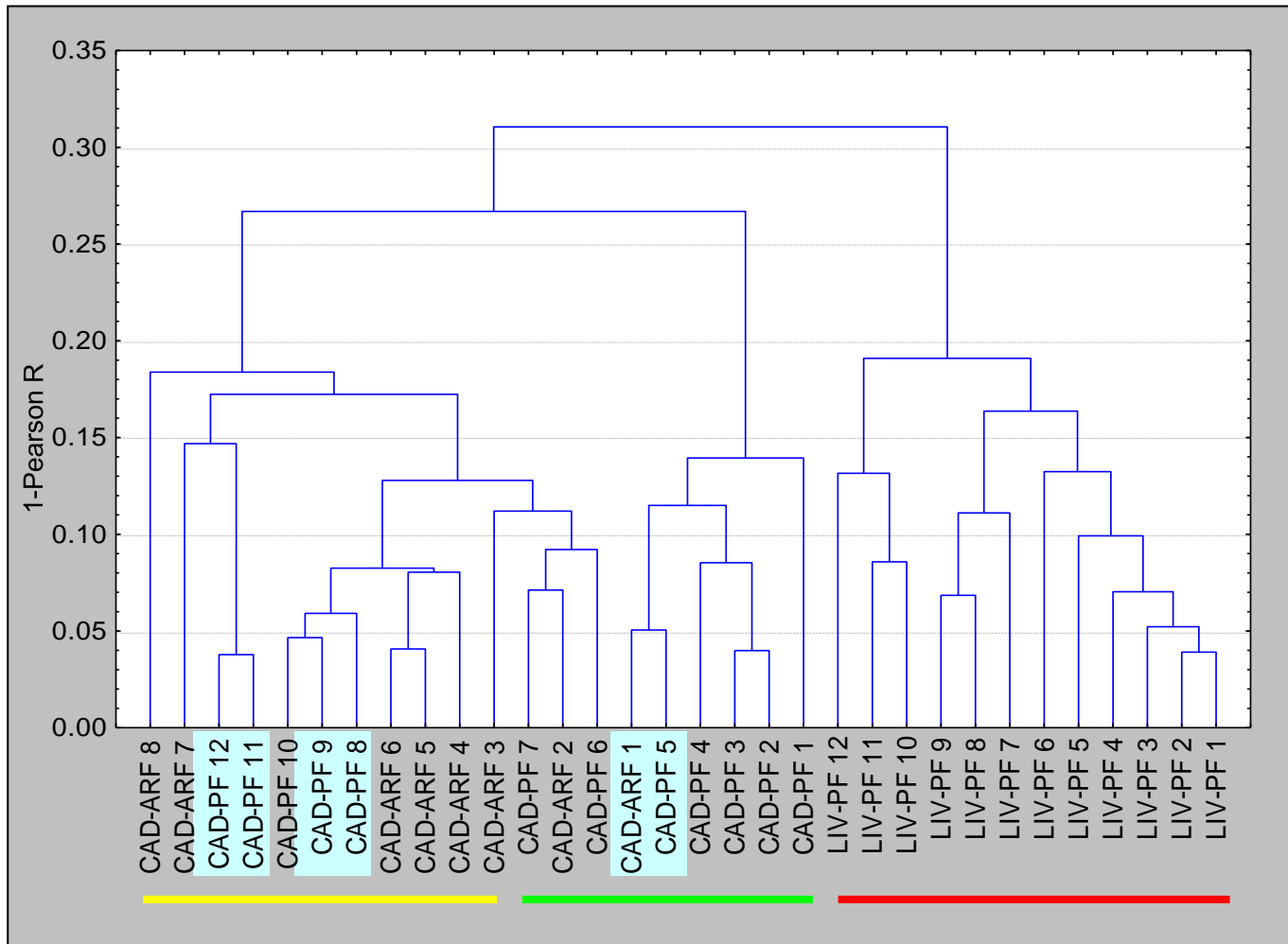
Glomeruli isolated from live and deceased donor kidney biopsies



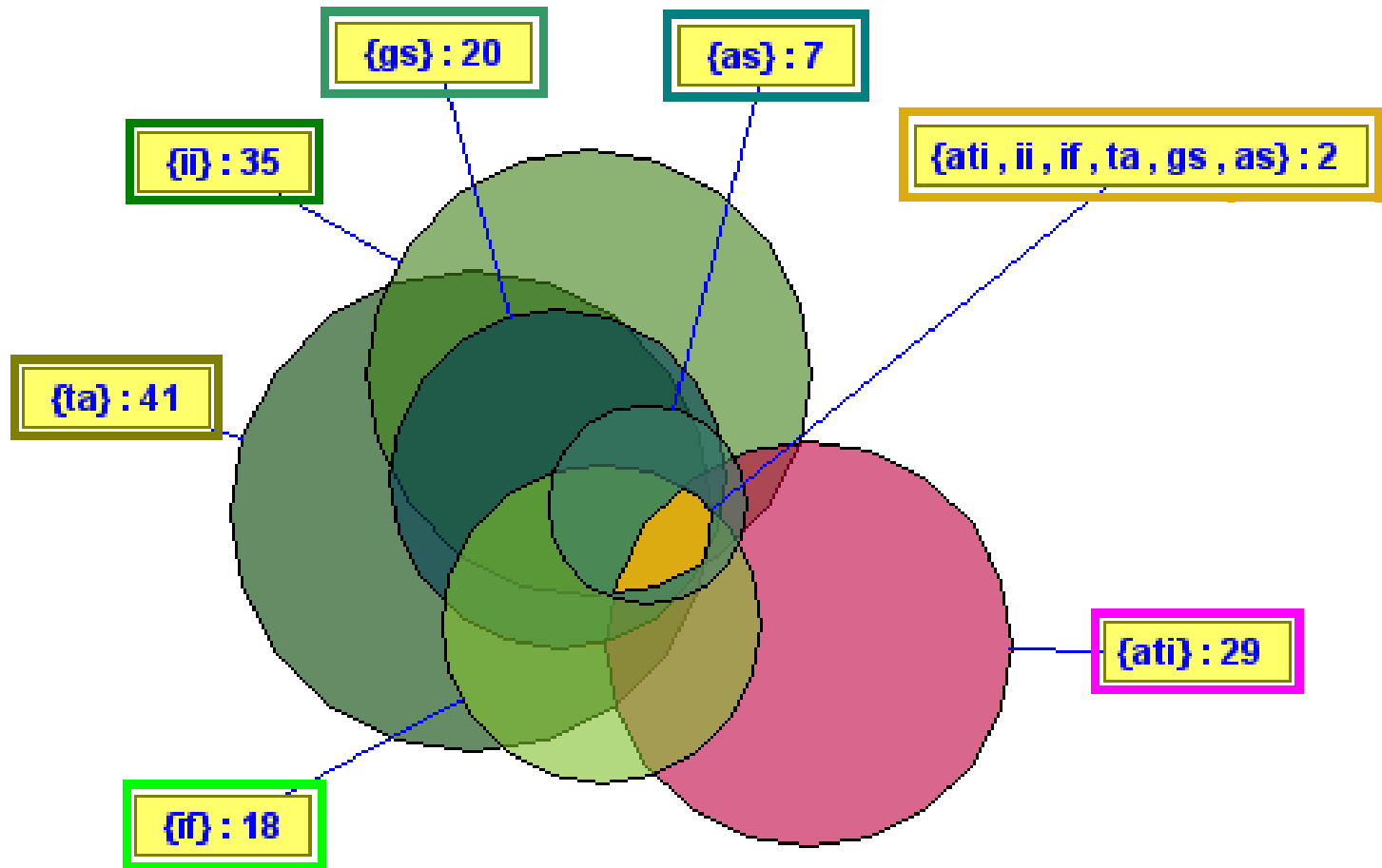
Transcripts are specific for deceased and live donor kidney compartments



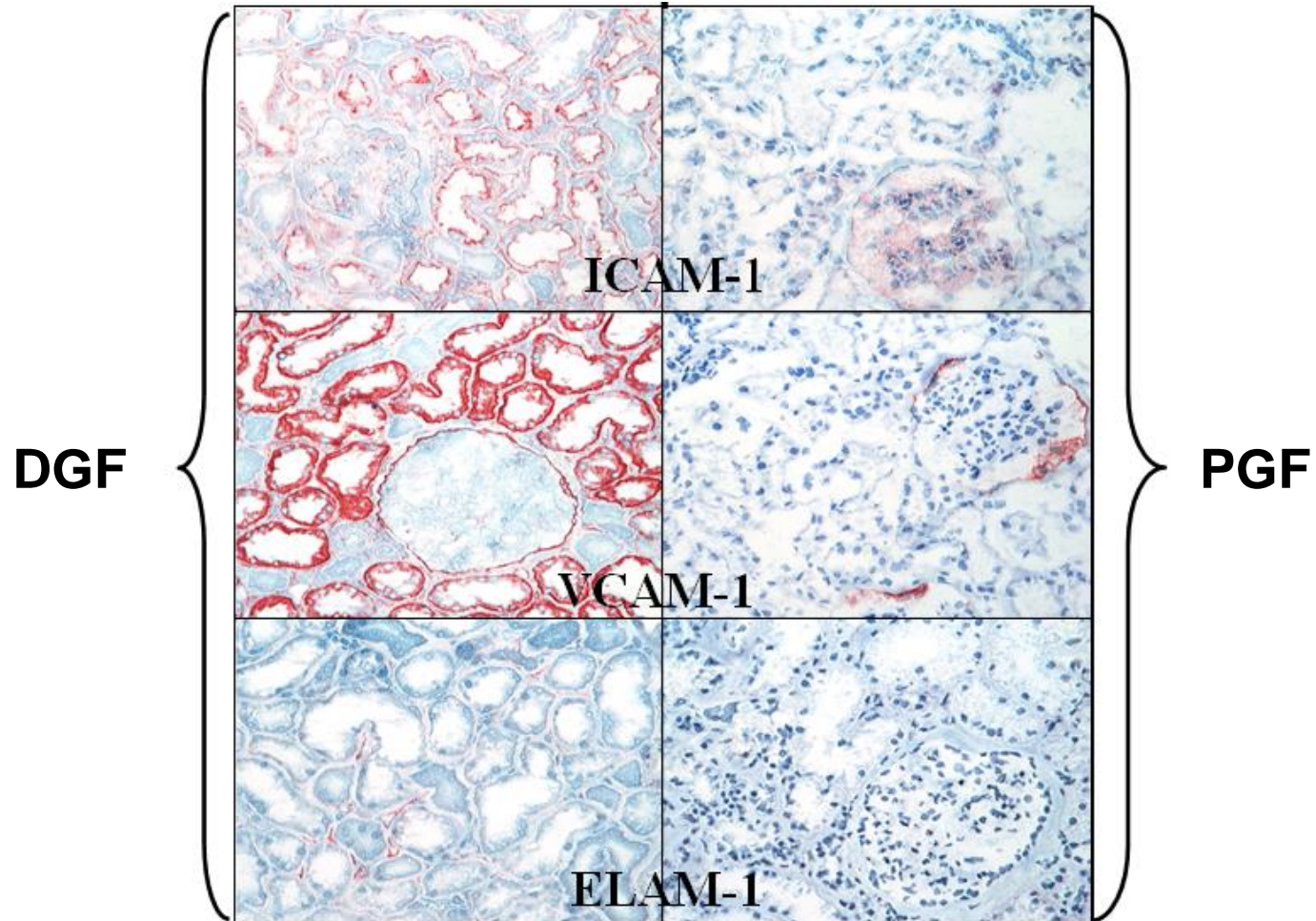
Unsupervised Cluster Analysis of Donor Kidney Biopsies



Histogenomics of deceased donor kidneys (n=82)

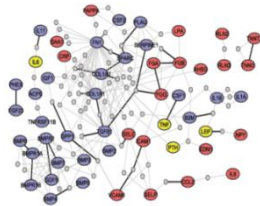


Inflammation in Deceased Donor Kidney Biopsies



Systems Biology and omicsNET

Physical Interactions

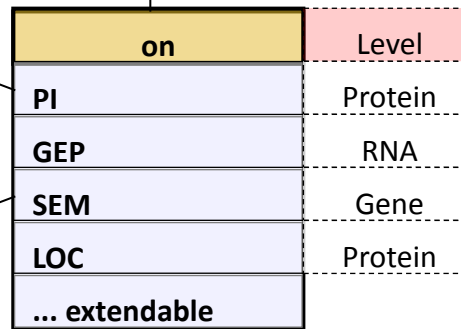


SEMantic Annotation Terms

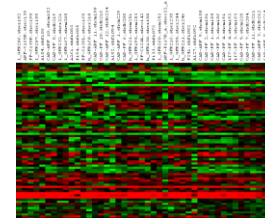
```

@all: all (170960) *
@GO:0001510: biological_process (130996)
@GO:0005675: cellular_component (107781) *
@GO:0005623: cell (71924)
@GO:0044864: cell part (17176)
@GO:0031975: envelope (2776)
@GO:0031012: extracellular matrix (579)
@GO:0044420: extracellular matrix part (200)
@GO:0005976: extracellular region (6249)
@GO:0044421: extracellular region part (4742)
@GO:0031974: membrane-enclosed lumen (4363)
@GO:0043226: organelle (46672)
@GO:0044422: organelle part (10269)
@GO:0043234: protein complex (13142)
@GO:0046202: synapse (265)
@GO:0044456: synapse part (101)
@GO:0019012: viron (151)
@GO:0044423: viron part (121)
@GO:0003674: molecular_function (126783)
@biomole_biological_process: biomole_biological_process (0)
@biomole_cellular_component: biomole_cellular_component (0)
@biomole_molecular_function: biomole_molecular_function (1)
    
```

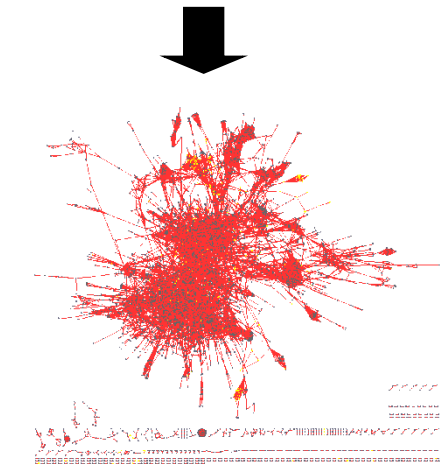
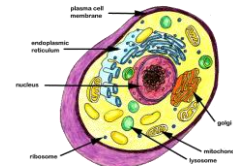
Label



Gene Expression Patterns

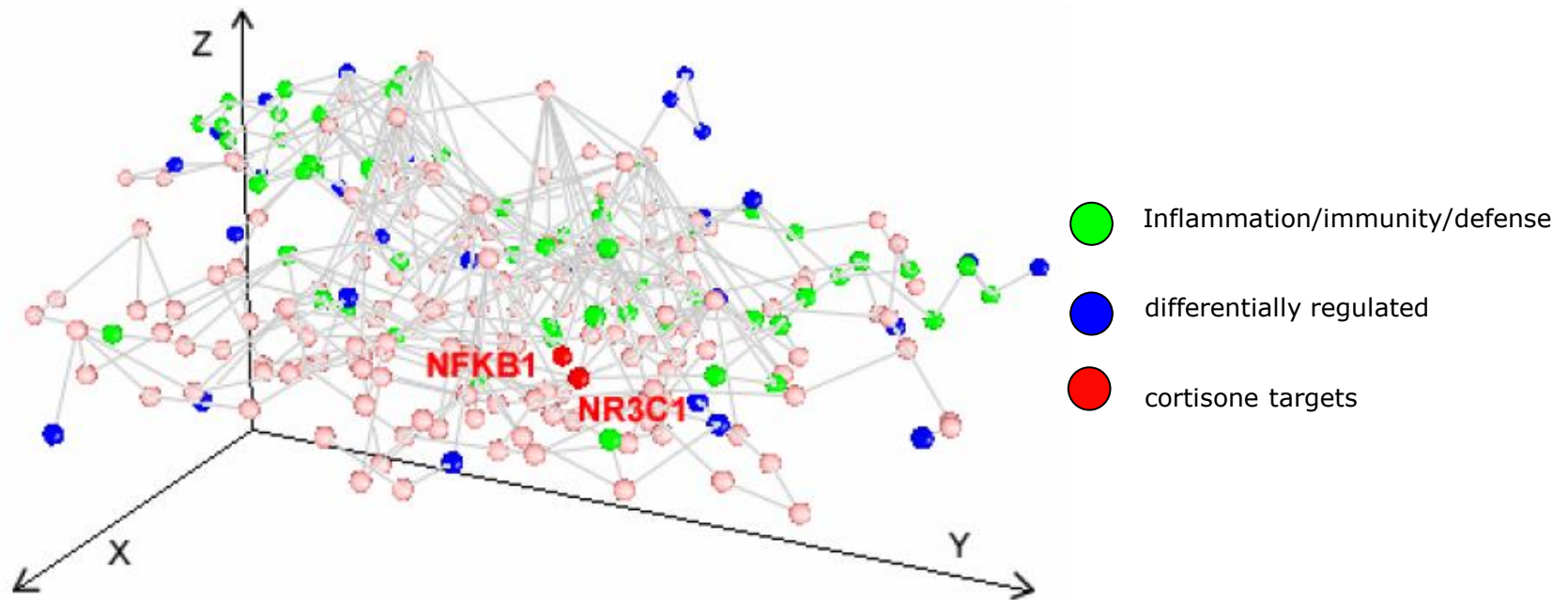


Subcellular Localization



omicsNET – Example – IRI/ARF

Inflammation networks in cadaveric donor organs

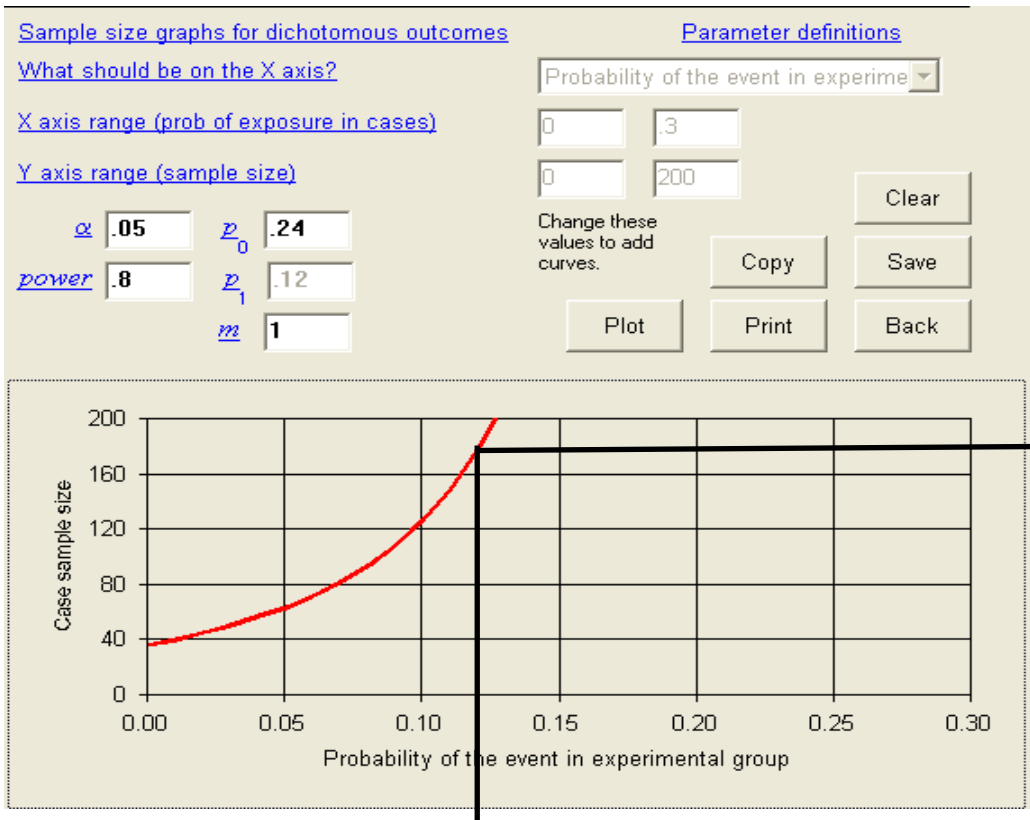


A Multicenter Double-Blinded RCT of Deceased Organ Donor Pre-Treatment with Corticosteroids for the Prevention of Postischemic Acute Renal Allograft Failure

Current Controlled Trials Registration #:
ISRCTN78828338

Sponsor: FWF P15679, €350k 3yrs

Sample size estimation



Event rate
control group = 24%

Computed sample
size to half event rate :

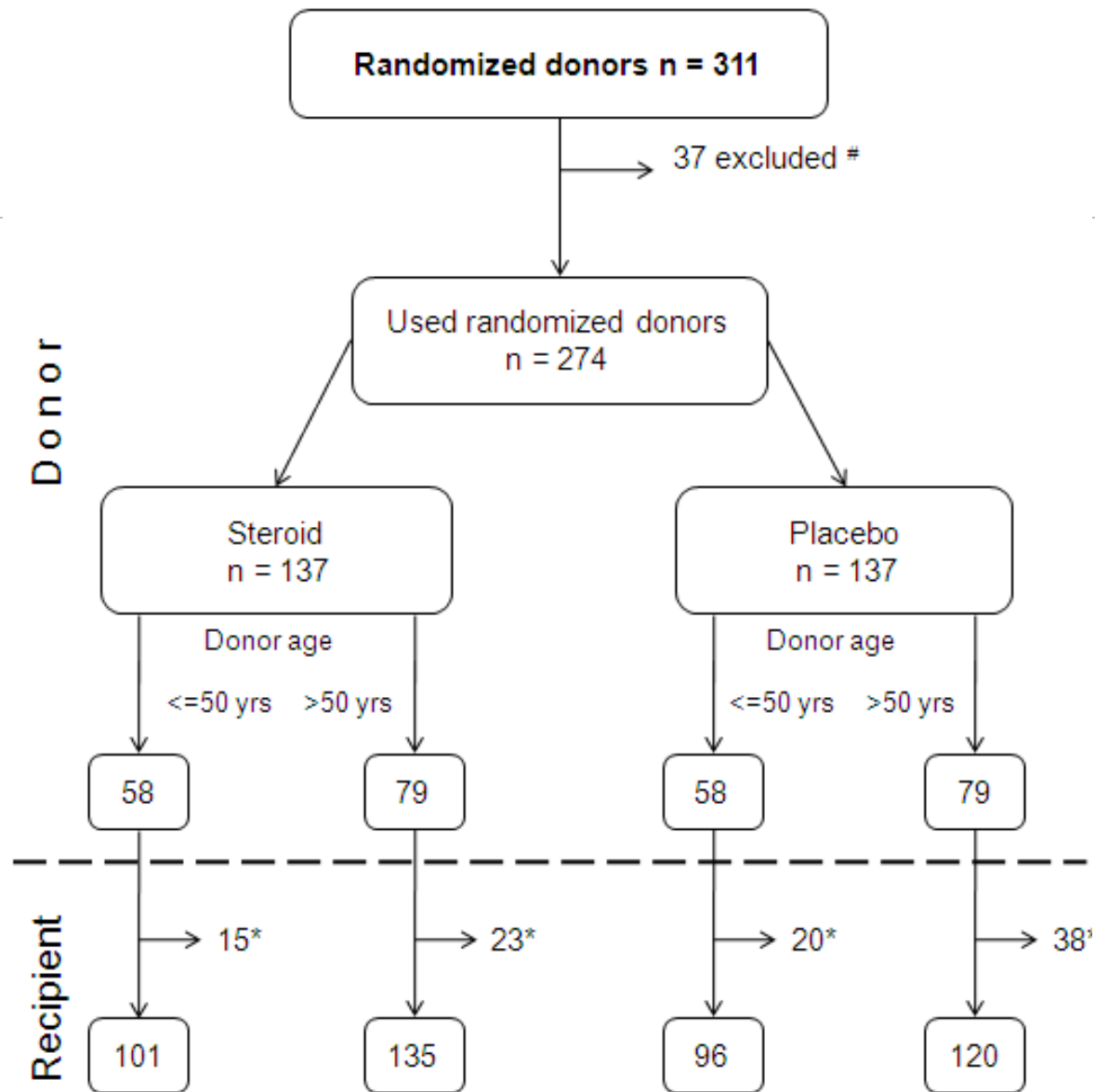
176

Donor sample size

$$N^* = N(1/1 - LFU/NU)^2 =$$

$$= 176(1/.92)^2 = 207$$

12%



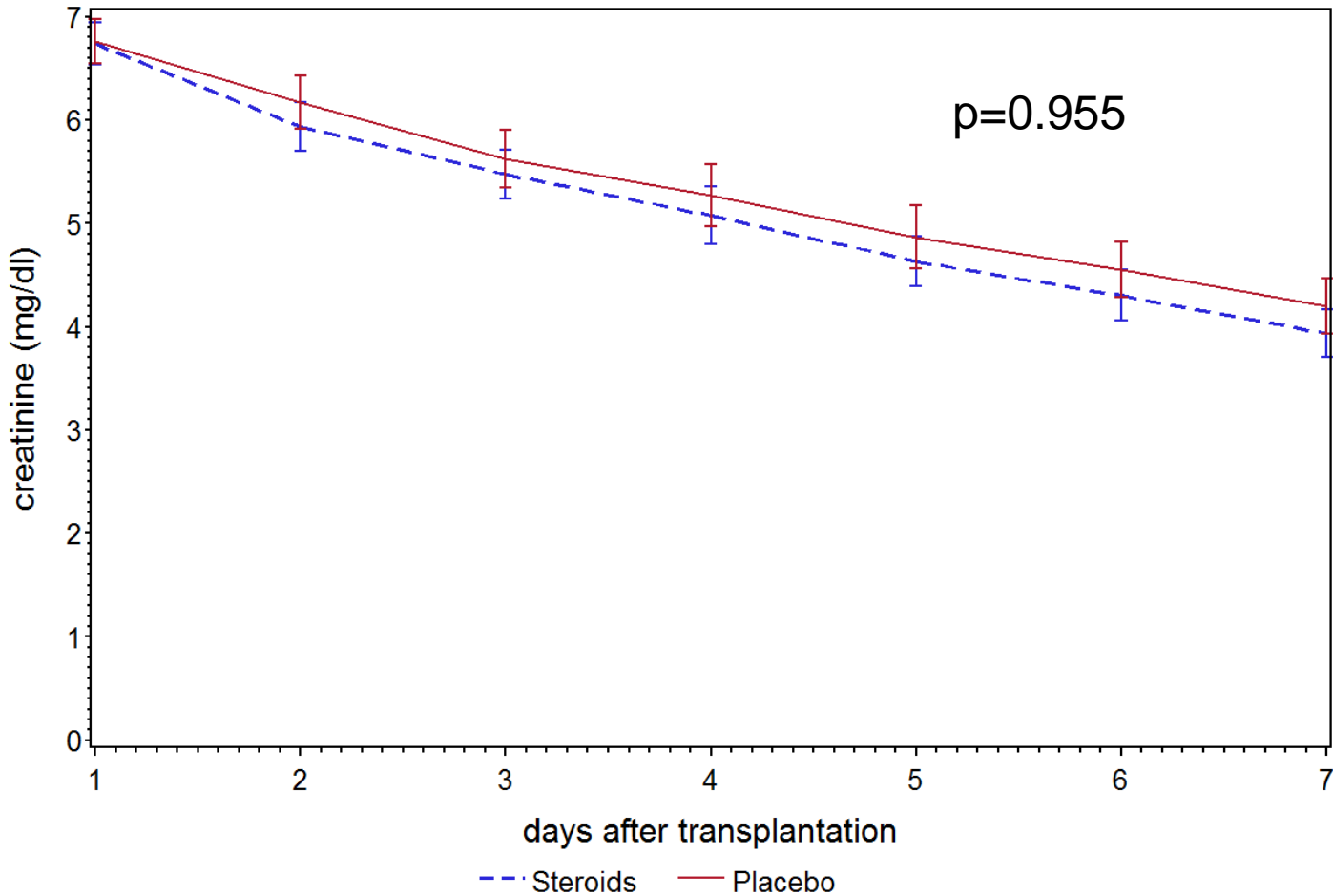
Demography of the 274 donors & 458 recipients

Variable	Steroid	Placebo	p-value
Number of donors	137	137	na
Donor age (years)	47.1 (15.1)	48.5 (14.0)	0.452
Donor sex (f/m)	62/74	57/76	0.652
Last creatinine of donor (mg/dl)	0.89 (0.29)	0.90 (0.39)	0.840
Vasopressors used (n/y)	22/114	12/121	0.078
Multiorgan donors (n/y)	106/30	94/39	0.173
Number of recipients	239	219	na
Recipient age (years)	49.6 (14.4)	49.2 (13.9)	0.753
Recipient sex (f/m)	76/162	77/142	0.465
Transplant number (1/2/3/4/5)	204/23/8/2/1	194/21/4/0/0	0.505*
Cold ischemic time (hours)	16.9 (13.4)	16.9 (15.2)	0.995
PRA latest (%)	6 (17)	4 (12)	0.172
Sum of HLA mismatches	3 (1)	3 (1)	0.379
Immunosuppression (CNI/else)	223/15	205/14	0.969
Induction therapy (none/antiCD25/ATG)	143/83/12	126/86/7	0.437

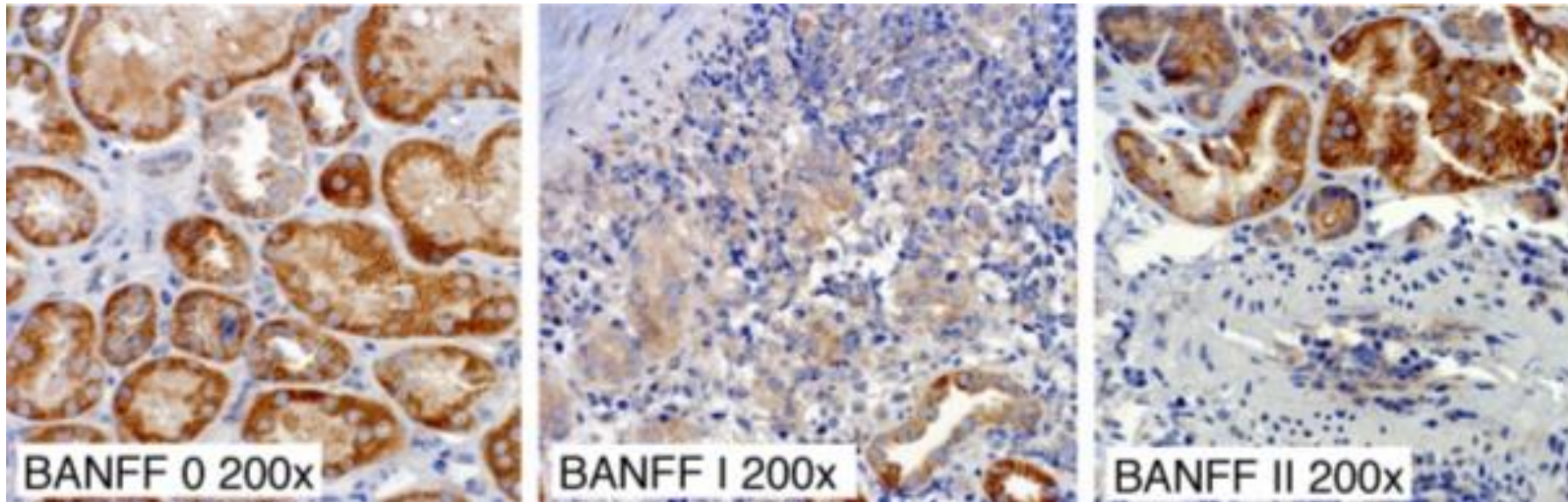
Primary study end point - DGF

	Steroids	Placebo	p-value
% Pts requiring dialysis during the first 7 days (0/1/>1)	65/13/22	63/12/25	0.700
Number of dialysis during the first 7 days (0/1/2/3/4/5)	154/32/18/28/2/4	137/27/27/18/8/2	0.115

Creatinine Trajectories



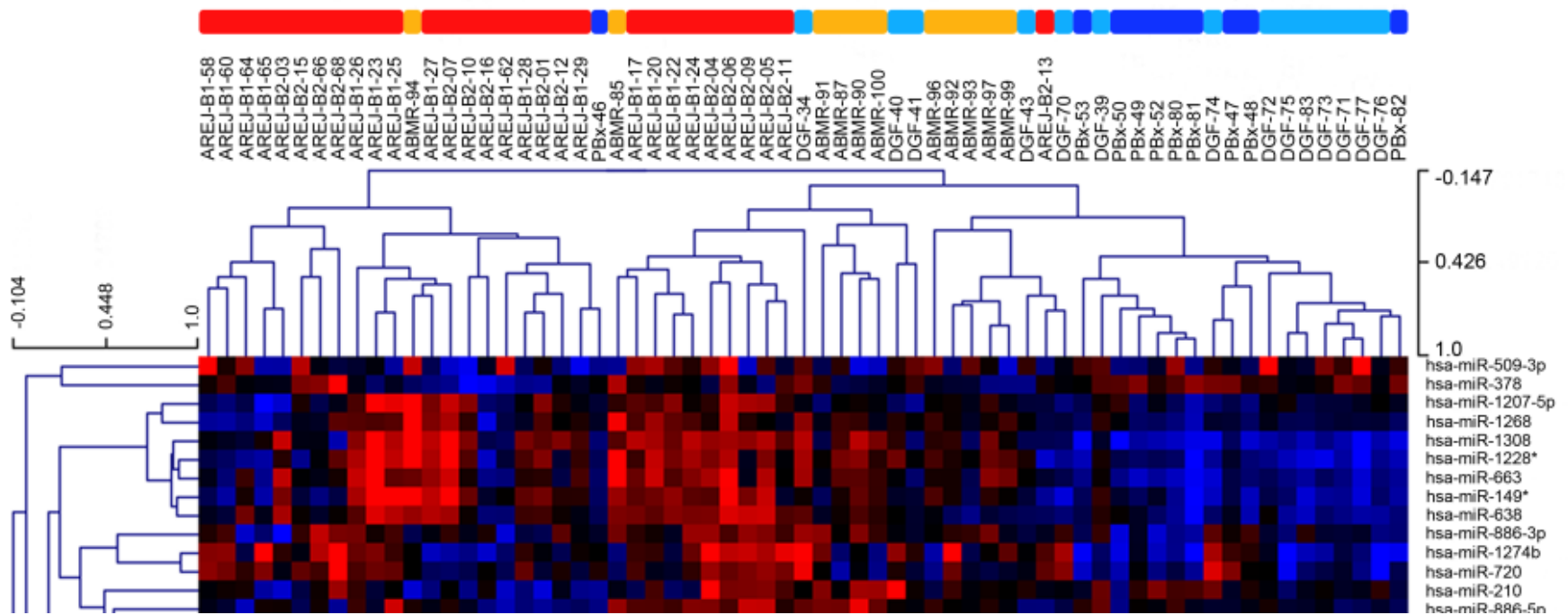
miRNA regulation of kidney injury (FWF P21436 - €250k 2010-2012)



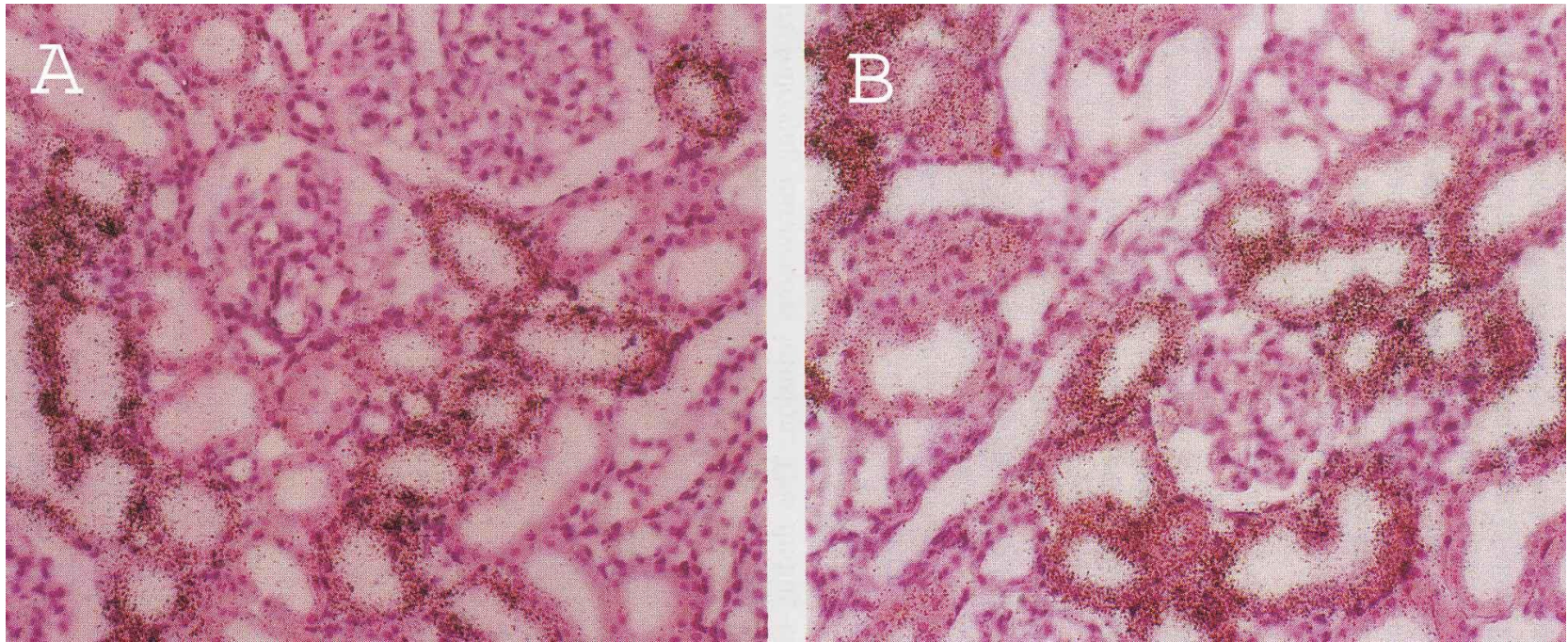
DICER

code for a complex cellular phenotype by silencing of target mRNAs

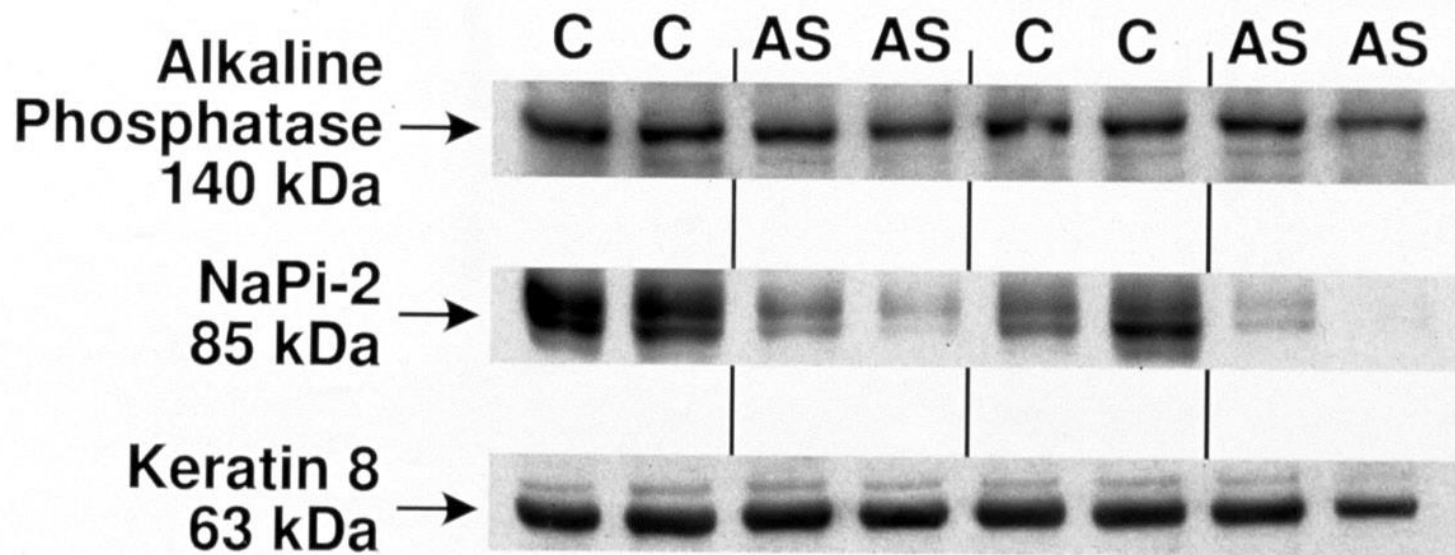
miRNA Profiles discriminate post-transplant events



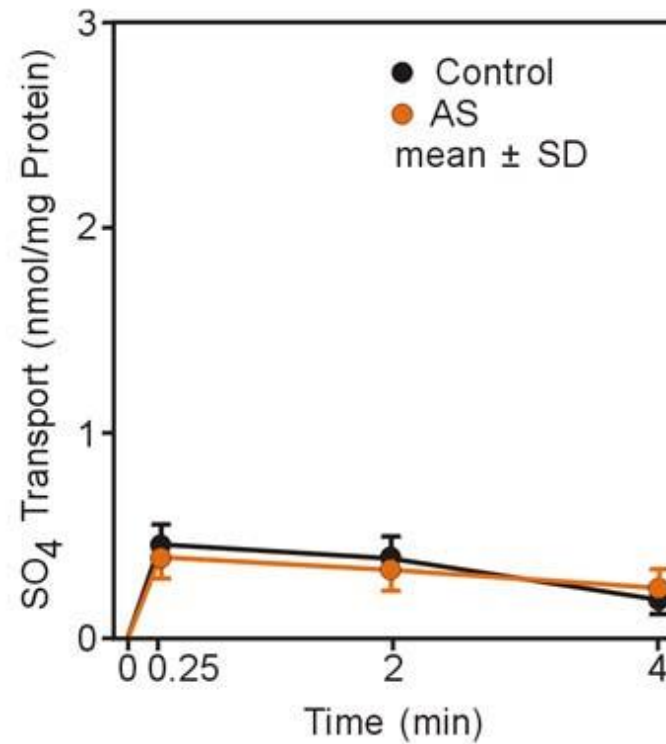
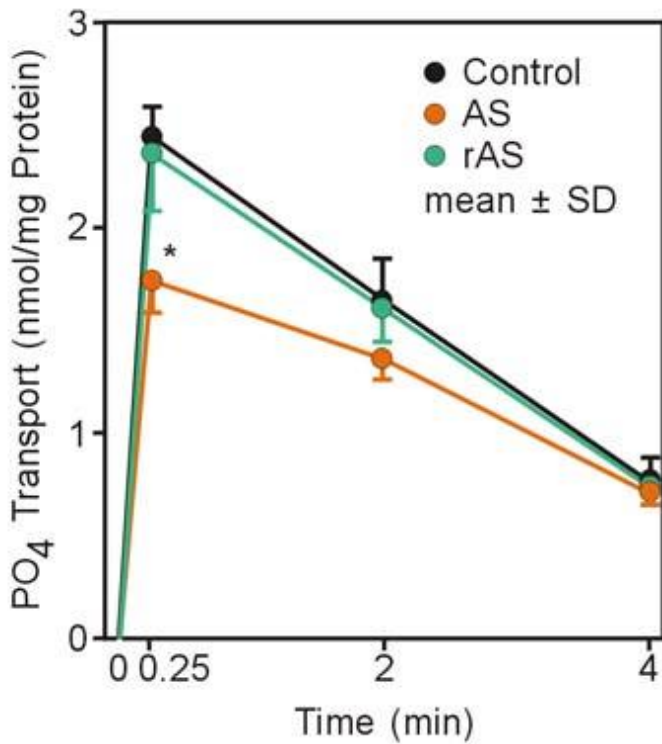
Antisense DNA/siRNA



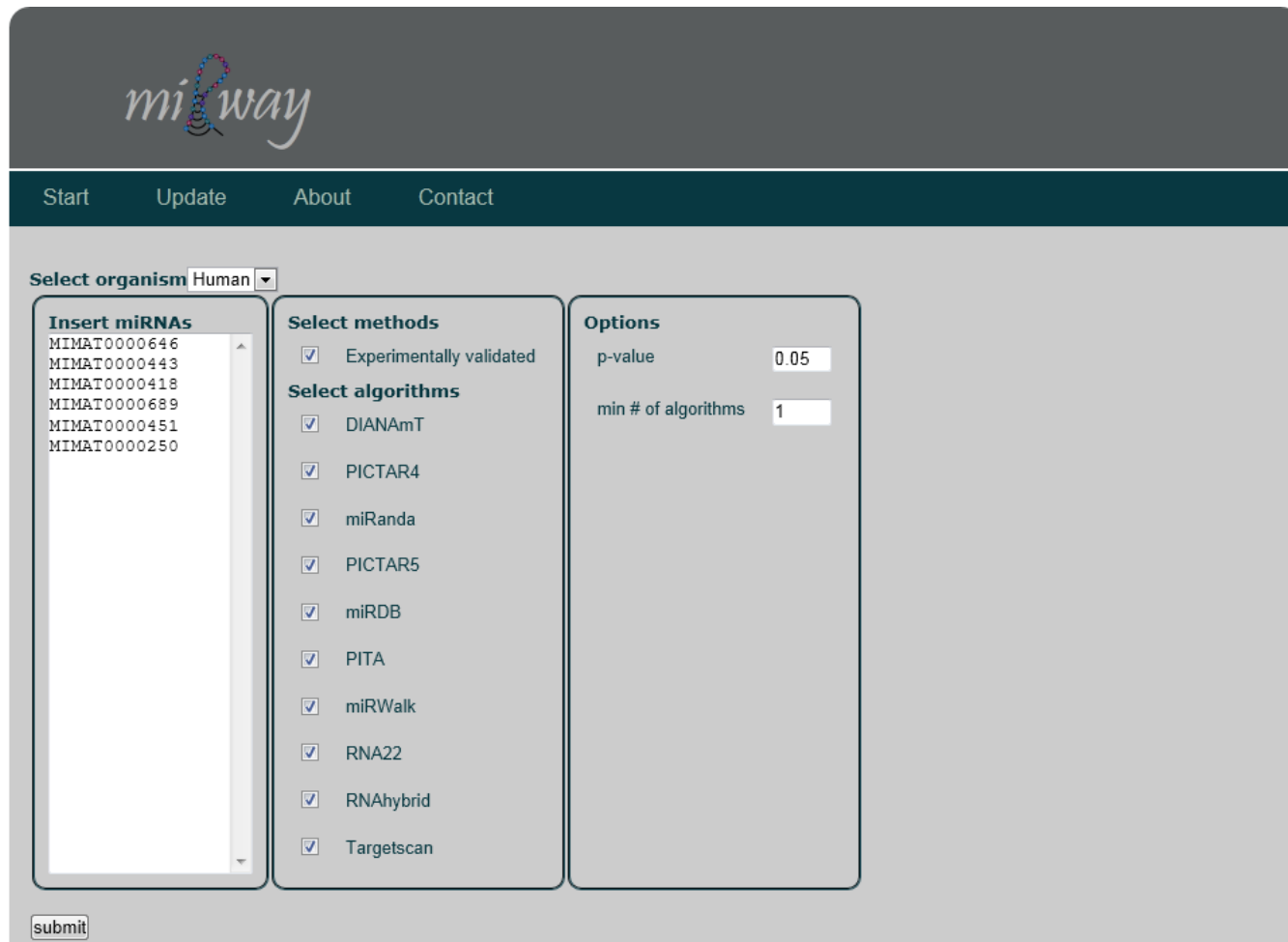
Antisense DNA/siRNA



Antisense DNA/siRNA



miRNA prediction tool



The screenshot shows the miRway web interface. At the top, there is a navigation bar with links for 'Start', 'Update', 'About', and 'Contact'. Below this, the 'miRway' logo is displayed. The main content area is divided into three sections:

- Select organism:** A dropdown menu currently set to 'Human'.
- Insert miRNAs:** A text input field containing a list of miRNA IDs: MIMAT0000646, MIMAT0000443, MIMAT0000418, MIMAT0000689, MIMAT0000451, and MIMAT0000250.
- Select methods:** A section with a checkbox for 'Experimentally validated'.
- Select algorithms:** A list of prediction algorithms, each with a checked checkbox: DIANA-T, PICTAR4, miRanda, PICTAR5, miRDB, PITA, miRWalk, RNA22, RNAhybrid, and Targetscan.
- Options:** Two input fields: 'p-value' set to 0.05 and 'min # of algorithms' set to 1.

A 'submit' button is located at the bottom left of the form area.

Acknowledgments



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BWH

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