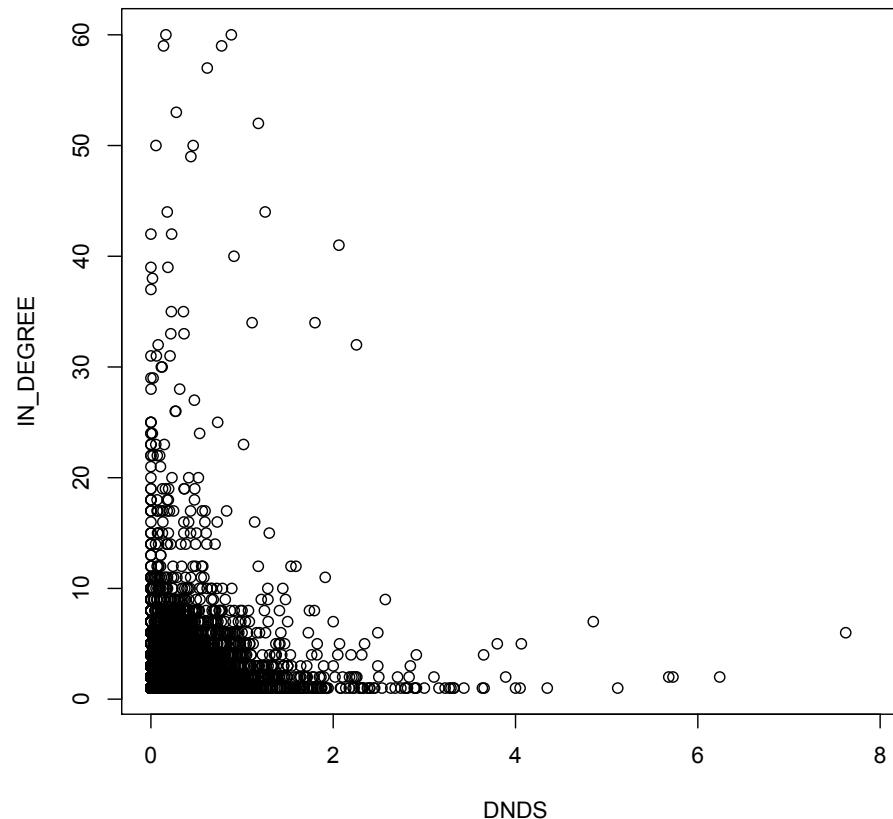


# Selection on TFs and target genes

Ekta Khurana, 29 Jul 2011

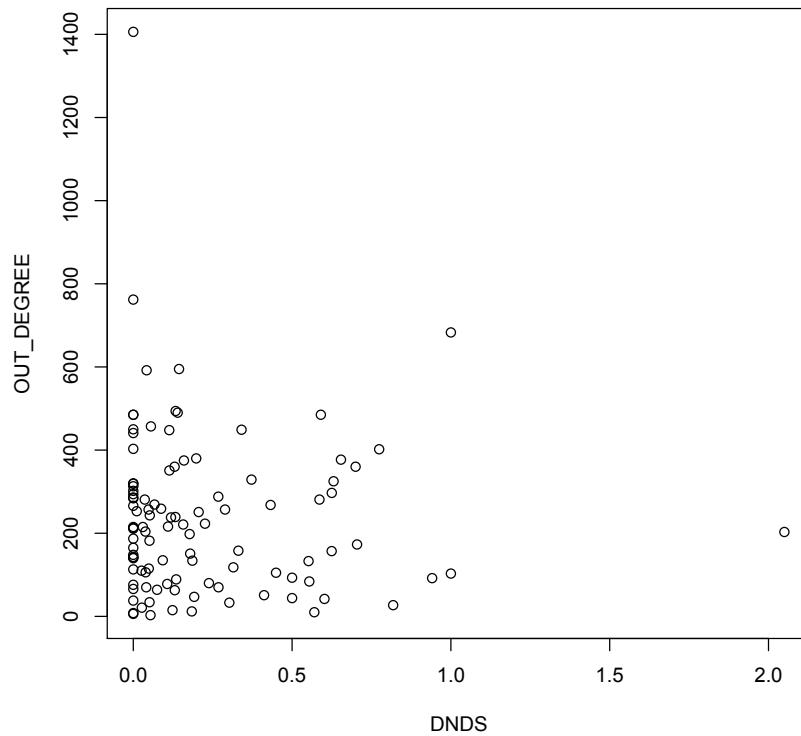
Targets with more regulators are under higher negative selection (chimp-human comparison)



TFs and Target genes included (in-degree > 0)

Spearman corr coeff=-0.05, p value = 7.6 e-06

# TFs and selection (chimp-human comparison)



**Out-degree > 0**

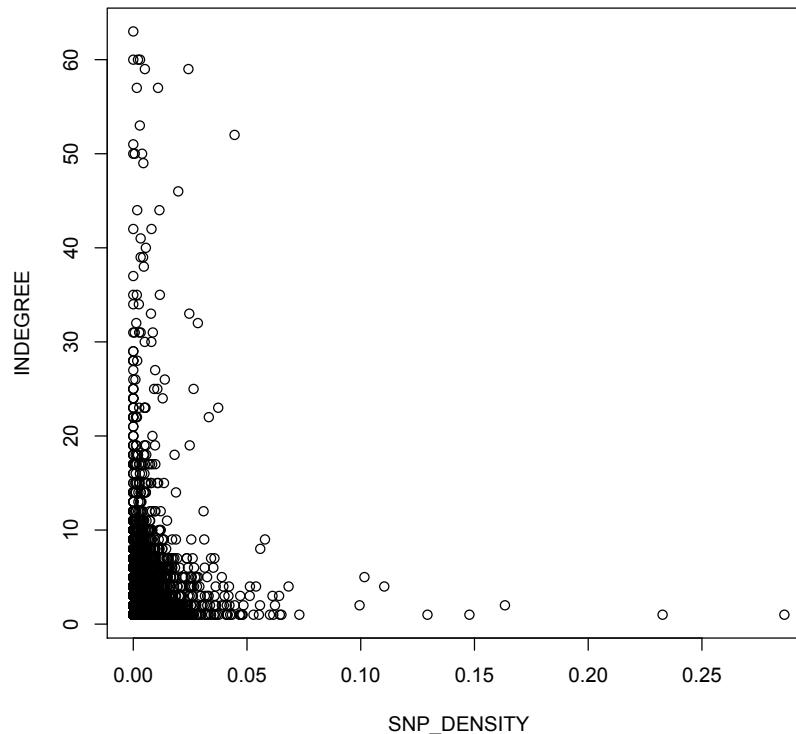
**Spearman corr coeff=-0.08, p value = 0.39**

# TFs contributing most to anti-correlation

TF name	dN/dS	Out-degree
PRDM1	0.818	27
CTCF	0	1406
RAD21	0	762
FAM48A	0.57	10
NRF1	0	485
GABPA	0	485
MEF2C	0.60	42
YY1	0	450
CCNT2	0.94	92
SP1	0	441

Targets with more regulators are under higher negative selection (within humans comparison)

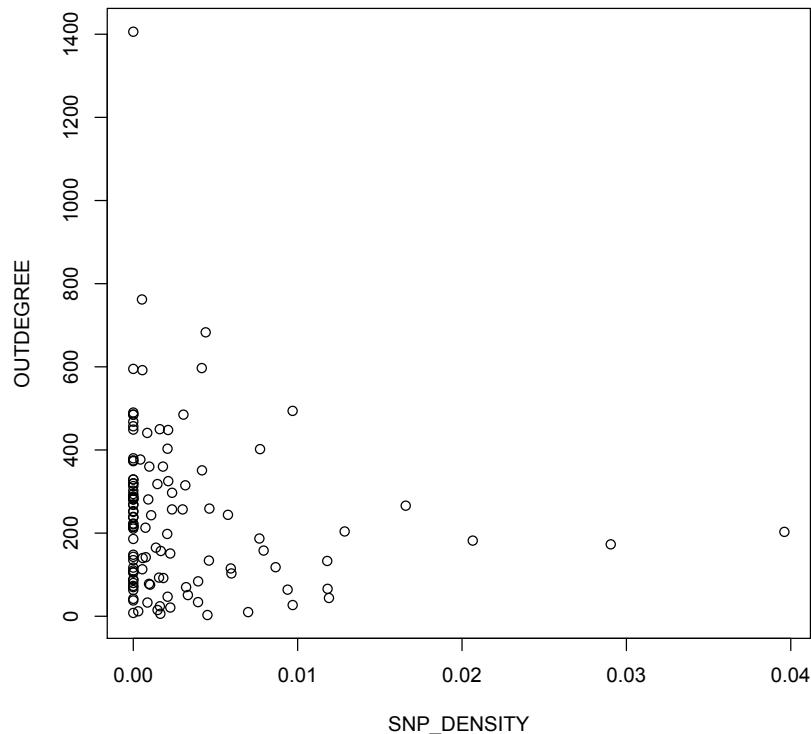
Non-synonymous SNPs from 1000 Genomes pilot project



**In-degree > 0**

Spearman corr coeff = - 0.05, pvalue=3.4e-05

TFs with more targets are under higher negative selection (within humans comparison)



**Out-degree > 0**

Spearman corr coeff = -0.19, pvalue=0.03

# TFs contributing most to anti-correlation

TF name	snp_Density	Out-degree
PRDM1	0.009	27
XRCC4	0.012	44
FAM48A	0.007	10
WRNIP1	0.005	3
ESRRA	0.012	66
CTCF	0	1406
SIRT6	0.009	64
TAF1	0	595
FOS	0	490
GABPA	0	485

# TFs contributing most to anti-correlation

For dN/dS	For SNP density
<b>PRDM1</b>	<b>PRDM1</b>
<b>CTCF</b>	<b>XRCC4</b>
<b>RAD21</b>	<b>FAM48A</b>
<b>FAM48A</b>	<b>WRNIP1</b>
<b>NRF1</b>	<b>ESRRA</b>
<b>GABPA</b>	<b>CTCF</b>
<b>MEF2C</b>	<b>SIRT6</b>
<b>YY1</b>	<b>TAF1</b>
<b>CCNT2</b>	<b>FOS</b>
<b>SP1</b>	<b>GABPA</b>