

Study Name	Public id	Bioset Name	Platform	Correlation/Direction	Common Genes	P-Value	Sample Count
Study_11:HEK293 cells DMSO treated cytoplasmic and pol	GSE45237	HEK293 cells DMSO treated - polyribosome asso	llumina iGenome UCSC, hg1	-	2057	1.60E-73	2
Study_22:Blood samples from healthy individuals; effects of	GSE53655	Blood samples from healthy female individuals - g	llumina iGenome UCSC, hg1	-	1293	1.10E-51	6
Study_22:Blood samples from healthy individuals; effects of	GSE53655	Blood samples from healthy male individuals - glo	llumina iGenome UCSC, hg1	-	974	5.60E-41	6
Study_22:Blood samples from healthy individuals; effects of	GSE53655	Whole blood samples non-globin depleted, from h	llumina iGenome UCSC, hg1	-	18	0.4791	6
Study_24:Hepatosplenic T cell lymphomas and peripheral b	GSE55716	Hepatosplenic T-cell lymphoma (HSTL) samples f	llumina iGenome UCSC, hg1	+	660	1.50E-50	3
Study_24:Hepatosplenic T cell lymphomas and peripheral b	GSE55716	Peripheral T-cell lymphoma (PTCL-NOS) sample	llumina iGenome UCSC, hg1	+	120	1.00E-21	3
Study_24:Hepatosplenic T cell lymphomas and peripheral b	GSE55716	Peripheral T-cell lymphoma samples - hepato	sple lllumina iGenome UCSC, hg1	+	293	5.10E-13	3
Study_24:Hepatosplenic T cell lymphomas and peripheral b	GSE55716	Hepatosplenic T-cell lymphoma (HTSL) DERL2 c	llumina iGenome UCSC, hg1	-	496	0.0455	1
Study_25:Chronic myeloid leukemia K562 cells; HuR (ELAV	GSE61238	Chronic myelogenous leukemia K562 cells - HuR	llumina iGenome UCSC, hg1	+	1633	3.40E-50	3
Study_31:Blood tissues of NLRC4-MAS and NOMID patient	GSE57253	Blood of a NLRC4-macrophage activation syndrom	llumina iGenome UCSC, hg1	-	1321	6.10E-47	3
Study_31:Blood tissues of NLRC4-MAS and NOMID patient	GSE57253	Blood of neonatal onset multisystem inflamma	llumina iGenome UCSC, hg1	-	131	1.20E-10	7
Study_31:Blood tissues of NLRC4-MAS and NOMID patient	GSE57253	Blood of neonatal onset multisystem inflamma	llumina iGenome UCSC, hg1	-	148	0.042	7
Study_46:Breast cancer MCF7 cells with EIF4A1 siRNA knc	GSE58111	Breast cancer MCF7 cells with EIF4A1 siRNA 48f	llumina iGenome UCSC, hg1	-	1703	3.70E-39	4
Study_46:Breast cancer MCF7 cells with EIF4A1 siRNA knc	GSE58111	Breast cancer MCF7 cells with control siRNA 48h	llumina iGenome UCSC, hg1	-	1593	2.10E-33	4
Study_46:Breast cancer MCF7 cells with EIF4A1 siRNA knc	GSE58111	Breast cancer MCF7 cells with EIF4A1 siRNA 48f	llumina iGenome UCSC, hg1	-	1180	5.00E-08	4
Study_46:Breast cancer MCF7 cells with EIF4A1 siRNA knc	GSE58111	Breast cancer MCF7 cells with control siRNA 48h	llumina iGenome UCSC, hg1	-	1059	3.00E-05	4
Study_46:Breast cancer MCF7 cells with EIF4A1 siRNA knc	GSE58111	Breast cancer MCF7 cells total RNA - EIF4A1 siR	llumina iGenome UCSC, hg1	-	483	0.1433	4
Study_46:Breast cancer MCF7 cells with EIF4A1 siRNA knc	GSE58111	Breast cancer MCF7 cells polysomal RNA - EIF4A	llumina iGenome UCSC, hg1	-	321	0.2077	4
Study_51:Endometria with or without endometriosis - molec	GSE51981	Endometria (proliferative) - mildly severe endome	Affymetrix GeneChip Human	-	2495	7.60E-37	11
Study_51:Endometria with or without endometriosis - molec	GSE51981	Endometria (mid-secretory) - mildly severe endom	Affymetrix GeneChip Human	-	2060	1.70E-36	9
Study_51:Endometria with or without endometriosis - molec	GSE51981	Endometria (mid-secretory) without endometriosis	Affymetrix GeneChip Human	-	1776	4.60E-34	14
Study_51:Endometria with or without endometriosis - molec	GSE51981	Endometria (proliferative) with endometriosis - m	c Affymetrix GeneChip Human	+	1571	4.30E-30	17
Study_51:Endometria with or without endometriosis - molec	GSE51981	Endometria (proliferative) without endometriosis	- Affymetrix GeneChip Human	-	2346	1.80E-24	15
Study_51:Endometria with or without endometriosis - molec	GSE51981	Endometria (mid-secretory) with endometriosis - r	Affymetrix GeneChip Human	+	1631	1.50E-21	18
Study_51:Endometria with or without endometriosis - molec	GSE51981	Endometria (mid-secretory) - moderately severe e	Affymetrix GeneChip Human	-	974	3.10E-21	18
Study_51:Endometria with or without endometriosis - molec	GSE51981	Endometria (proliferative) - moderately severe en	c Affymetrix GeneChip Human	-	2098	3.30E-16	17
Study_51:Endometria with or without endometriosis - molec	GSE51981	Endometria (early secretory) with endometriosis	- Affymetrix GeneChip Human	-	238	0.0222	12
Study_51:Endometria with or without endometriosis - molec	GSE51981	Endometria (early secretory) - moderately severe	Affymetrix GeneChip Human	-	1507	0.0305	12
Study_51:Endometria with or without endometriosis - molec	GSE51981	Endometria (early secretory) - mildly severe endo	Affymetrix GeneChip Human	-	586	0.0327	6
Study_52:Adipose tissue changes associated with diet and//	GSE43471	Subcutaneous adipose tissue of obese women - e	llumina Human HT-12 (WG-t	-	404	8.00E-37	14
Study_52:Adipose tissue changes associated with diet and//	GSE43471	Subcutaneous adipose tissue of obese women - s	llumina Human HT-12 (WG-t	-	94	3.00E-07	8
Study_52:Adipose tissue changes associated with diet and//	GSE43471	Subcutaneous adipose tissue of obese women - c	llumina Human HT-12 (WG-t	+	71	0.0066	16
Study_52:Adipose tissue changes associated with diet and//	GSE43471	Subcutaneous adipose tissue of obese women - c	llumina Human HT-12 (WG-t	-	80	0.0223	9
Study_52:Adipose tissue changes associated with diet and//	GSE43471	Subcutaneous adipose tissue of obese women - c	llumina Human HT-12 (WG-t	-	91	0.069	9
Study_52:Adipose tissue changes associated with diet and//	GSE43471	Subcutaneous adipose tissue of obese women - c	llumina Human HT-12 (WG-t	+	65	0.149	16
Study_52:Adipose tissue changes associated with diet and//	GSE43471	Subcutaneous adipose tissue of obese women - e	llumina Human HT-12 (WG-t	-	33	0.1601	14
Study_53:Motor neurons derived from iPSC cells from amyotr	GSE54409	Motor neurons derived from iPSCS heterozygous	llumina iGenome UCSC, hg1	+	912	1.90E-36	2
Study_56:Perturbation of the translome by the anti-diabeti	GSE36847	MCF7 breast cancer cells 1uM PP242 treated 12f	Affymetrix GeneChip Human	+	2473	7.40E-36	4
Study_56:Perturbation of the translome by the anti-diabeti	GSE36847	MCF7 breast cancer cells 10mM metformin treate	Affymetrix GeneChip Human	+	2374	2.00E-34	4
Study_56:Perturbation of the translome by the anti-diabeti	GSE36847	MCF7 breast cancer cells vehicle control 12hr - p	c Affymetrix GeneChip Human	+	2288	4.20E-28	4
Study_56:Perturbation of the translome by the anti-diabeti	GSE36847	MCF7 breast cancer cells 100mM rapamycin treat	Affymetrix GeneChip Human	+	2175	6.40E-27	4
Study_56:Perturbation of the translome by the anti-diabeti	GSE36847	MCF7 breast cancer cytosolic RNA - 1uM PP242	Affymetrix GeneChip Human	-	420	6.80E-06	4
Study_56:Perturbation of the translome by the anti-diabeti	GSE36847	MCF7 breast cancer polysome RNA - 100mM rap	Affymetrix GeneChip Human	-	367	0.0006	4
Study_56:Perturbation of the translome by the anti-diabeti	GSE36847	MCF7 breast cancer polysome RNA - 10mM metf	Affymetrix GeneChip Human	+	510	0.0012	4
Study_56:Perturbation of the translome by the anti-diabeti	GSE36847	MCF7 breast cancer cytosolic RNA - 10mM metfo	Affymetrix GeneChip Human	+	197	0.0136	4
Study_56:Perturbation of the translome by the anti-diabeti	GSE36847	MCF7 breast cancer polysome RNA - 1uM PP242	Affymetrix GeneChip Human	+	858	0.0241	4
Study_57:Bronchial epithelial gene expression in cells treat	E-TABM-127	Bronchial epithelial cells (no recovery) treated	1hr Affymetrix GeneChip Human	-	1168	8.50E-36	3
Study_57:Bronchial epithelial gene expression in cells treat	E-TABM-127	Bronchial epithelial cells treated 1hr with tobacco	Affymetrix GeneChip Human	-	882	5.80E-16	3
Study_57:Bronchial epithelial gene expression in cells treat	E-TABM-127	Bronchial epithelial cells treated 1hr with tobacco	Affymetrix GeneChip Human	-	974	5.10E-10	3
Study_57:Bronchial epithelial gene expression in cells treat	E-TABM-127	Bronchial epithelial cells (after 5hr recovery in cult	Affymetrix GeneChip Human	-	998	9.60E-09	3
Study_57:Bronchial epithelial gene expression in cells treat	E-TABM-127	Bronchial epithelial cells treated 1hr with tobacco	Affymetrix GeneChip Human	-	750	3.10E-06	3