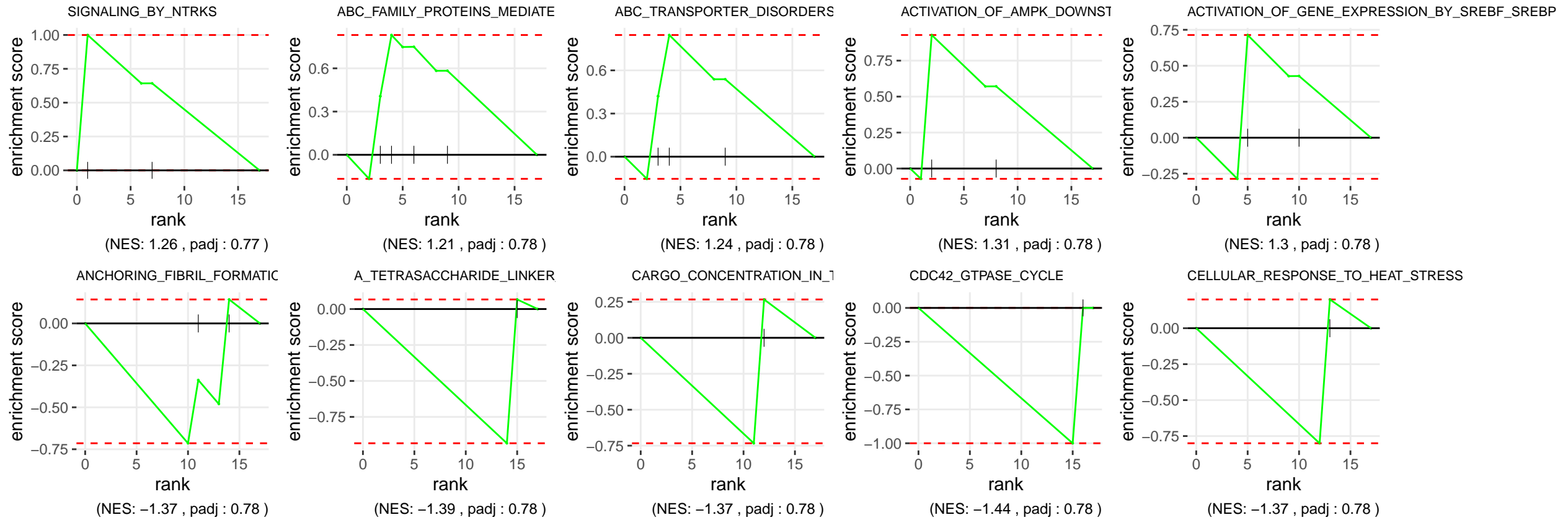


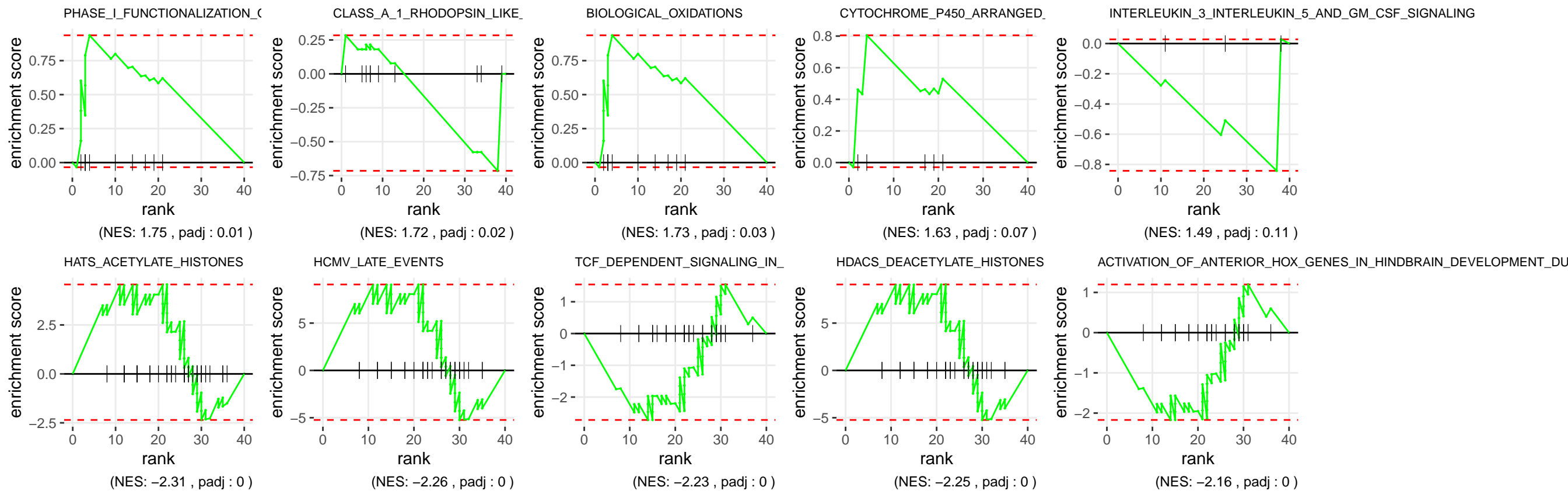
D2 ECs : Top enriched Pathways (GSEA), Old vs Young

Pathway	Gene ranks	NES	pval	padj
REACTOME_SIGNALING_BY_NTRKS		1.26	1.5e-03	7.7e-01
REACTOME_ABC_FAMILY_PROTEINS_MEDIATED_TRANSPORT		1.21	1.7e-01	7.8e-01
REACTOME_ABC_TRANSPORTER_DISORDERS		1.24	1.6e-01	7.8e-01
REACTOME_ACTIVATION_OF_AMPK_DOWNSTREAM_OF_NMDARS		1.31	4.0e-02	7.8e-01
REACTOME_ACTIVATION_OF_GENE_EXPRESSION_BY_SREBF_SREBP		1.30	1.8e-01	7.8e-01
REACTOME_CELLULAR_RESPONSE_TO_HEAT_STRESS		-1.37	2.1e-01	7.8e-01
REACTOME_CDC42_GTPASE_CYCLE		-1.44	2.0e-01	7.8e-01
REACTOME_CARGO_CONCENTRATION_IN_THE_ER		-1.37	2.0e-01	7.8e-01
REACTOME_A_TETRASACCHARIDE_LINKER_SEQUENCE_IS_REQUIRED_FOR_GAG_SYNTHESIS		-1.39	1.1e-01	7.8e-01
REACTOME_ANCHORING_FIBRIL_FORMATION		-1.37	2.0e-01	7.8e-01



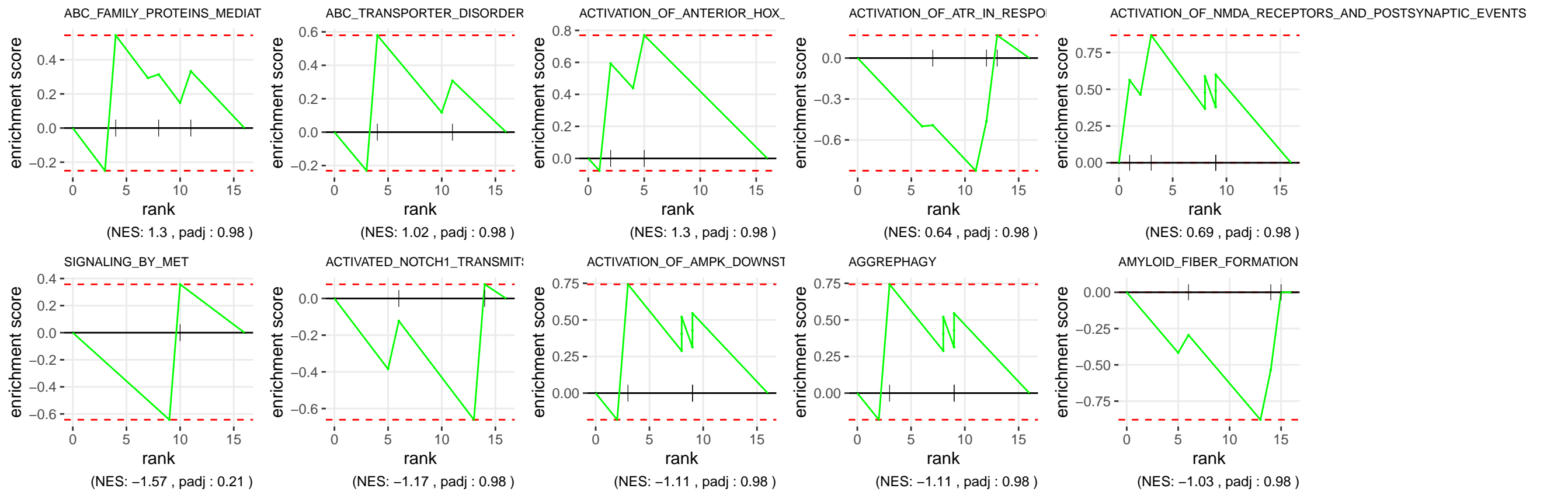
D2 FAPs : Top enriched Pathways (GSEA), Old vs Young

	Pathway	Gene ranks	NES	pval	padj
	REACTOME_PHASE_I_FUNCTIONALIZATION_OF_COMPOUNDS	1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40	1.75	1.6e-03	1.3e-02
	REACTOME_CLASS_A_1_RHODOPSIN_LIKE_RECEPTORS	1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40	1.72	2.8e-03	2.2e-02
	REACTOME_BIOLOGICAL_OXIDATIONS	1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40	1.73	4.6e-03	3.4e-02
	REACTOME_CYTOCHROME_P450_ARRANGED_BY_SUBSTRATE_TYPE	1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40	1.63	1.0e-02	6.7e-02
	REACTOME_INTERLEUKIN_3_INTERLEUKIN_5_AND_GM-CSF_SIGNALING	1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40	1.49	1.7e-02	1.1e-01
	REACTOME_ACTIVATION_OF_ANTERIOR_HOX_GENES_IN_HINDBRAIN_DEVELOPMENT_DURING_EARLY_EMBRYOGENESIS	1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40	-2.16	2.8e-06	1.8e-04
	REACTOME_HDACs_DEACETYLATE_HISTONES	1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40	-2.25	5.2e-07	7.4e-05
	REACTOME_TCF_DEPENDENT_SIGNALING_IN_RESPONSE_TO_WNT	1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40	-2.23	2.9e-07	5.5e-05
	REACTOME_HCMV_LATE_EVENTS	1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40	-2.26	1.2e-07	3.4e-05
	REACTOME_HATS_ACETYLATE_HISTONES	1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40	-2.31	8.5e-08	3.4e-05






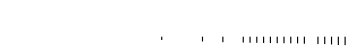






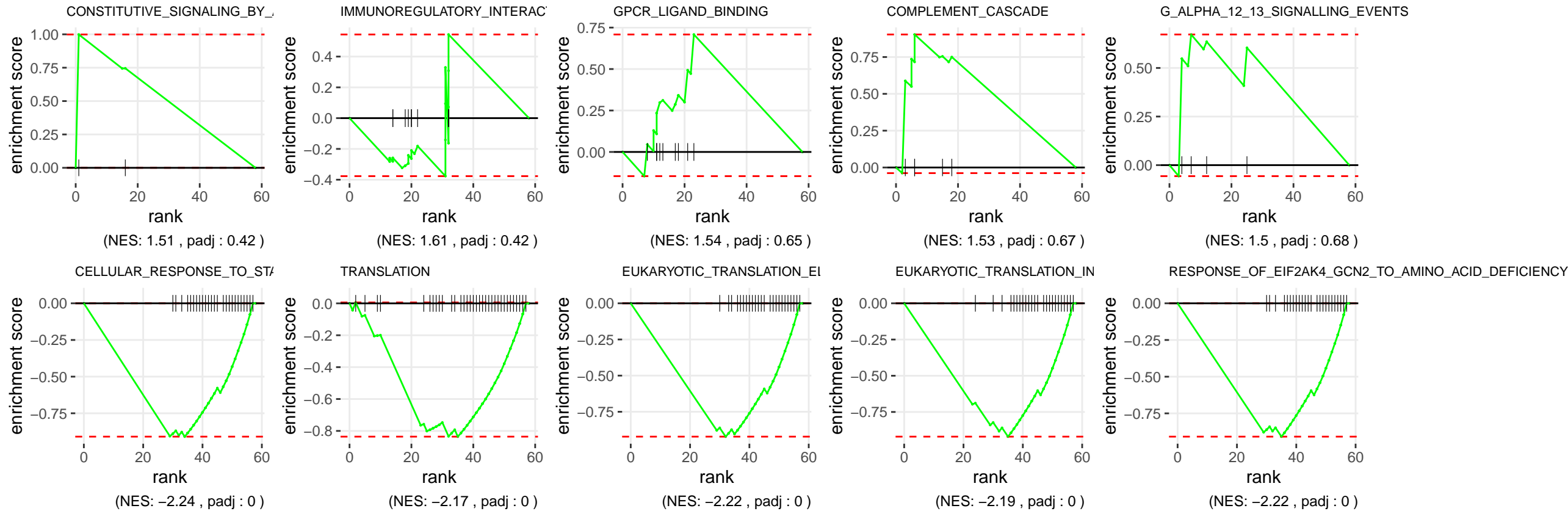
D2 M1 : Top enriched Pathways (GSEA), Old vs Young

	Pathway	Gene ranks	NES	pval	padj
	REACTOME_ABC_FAMILY_PROTEINS_MEDIATED_TRANSPORT		1.30	1.9e-01	9.8e-01
	REACTOME_ABC_TRANSPORTER_DISORDERS		1.02	4.8e-01	9.8e-01
	ACTIVATION_OF_ANTERIOR_HOX_GENES_IN_HINDBRAIN_DEVELOPMENT_DURING_EARLY_EMBRYOGENESIS		1.30	1.8e-01	9.8e-01
	REACTOME_ACTIVATION_OF_ATR_IN_RESPONSE_TO_REPLICATION_STRESS		0.64	8.0e-01	9.8e-01
	REACTOME_ACTIVATION_OF_NMDA_RECEPTORS_AND_POSTSYNAPTIC_EVENTS		0.69	7.7e-01	9.8e-01
	REACTOME_AMYLOID_FIBER_FORMATION		-1.03	4.2e-01	9.8e-01
	REACTOME_AGGREPHAGY		-1.11	3.5e-01	9.8e-01
	REACTOME_ACTIVATION_OF_AMPK_DOWNSTREAM_OF_NMDARS		-1.11	3.5e-01	9.8e-01
	REACTOME_ACTIVATED_NOTCH1_TRANSMITS_SIGNAL_TO_THE_NUCLEUS		-1.17	2.9e-01	9.8e-01
	REACTOME_SIGNALING_BY_MET		-1.57	4.0e-04	2.1e-01



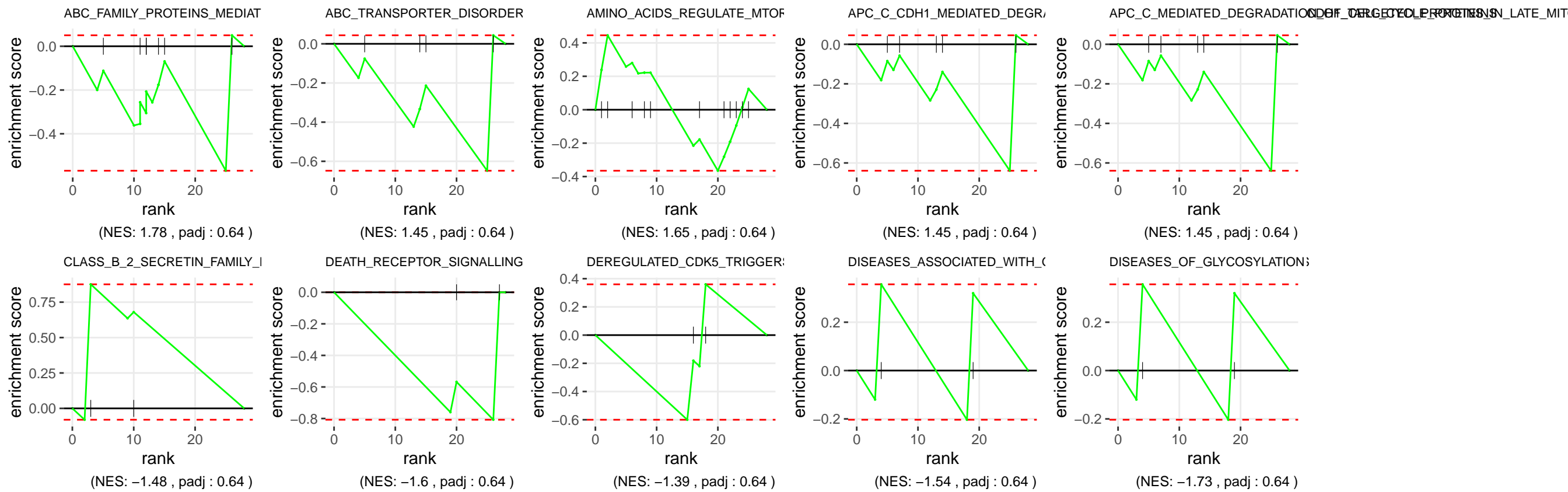
D2 M2 : Top enriched Pathways (GSEA), Old vs Young

	Pathway	Gene ranks	NES	pval	padj
	REACTOME_CONSTITUTIVE_SIGNALING_BY_ABERRANT_PI3K_IN_CANCER		1.51	1.4e-02	4.2e-01
	REACTOME_IMMUNOREGULATORY_INTERACTIONS_BETWEEN_A_LYMPHOID_AND_A_NON_LYMPHOID_CELL		1.61	1.5e-02	4.2e-01
	REACTOME_GPCR_LIGAND_BINDING		1.54	3.8e-02	6.5e-01
	REACTOME_COMPLEMENT_CASCADE		1.53	4.1e-02	6.7e-01
	REACTOME_G_ALPHA_12_13_SIGNALLING_EVENTS		1.50	4.6e-02	6.8e-01
	REACTOME_RESPONSE_OF_EIF2AK4_GC2N2_TO_AMINO_ACID_DEFICIENCY		-2.22	9.3e-06	9.0e-04
	REACTOME_EUKARYOTIC_TRANSLATION_INITIATION		-2.19	7.1e-06	9.0e-04
	REACTOME_EUKARYOTIC_TRANSLATION_ELONGATION		-2.22	9.3e-06	9.0e-04
	REACTOME_TRANSLATION		-2.17	2.1e-06	7.5e-04
	REACTOME_CELLULAR_RESPONSE_TO_STARVATION		-2.24	2.6e-06	7.5e-04
		0 10 20 30 40 50			













D2 Neutro : Top enriched Pathways (GSEA), Old vs Young

	Pathway	Gene ranks	NES	pval	padj
	REACTOME_ABC_FAMILY_PROTEINS_MEDIATED_TRANSPORT	0 5 10 15 20 25	1.78	1.3e-02	6.4e-01
	REACTOME_ABC_TRANSPORTER_DISORDERS	0 5 10 15 20 25	1.45	1.0e-01	6.4e-01
	REACTOME_AMINO_ACIDS_REGULATE_MTORC1	0 5 10 15 20 25	1.65	2.2e-02	6.4e-01
	DEGRADATION_OF_CDC20_AND_OTHER_APC_C_CDH1_TARGETED_PROTEINS_IN_LATE_MITOSIS_EARLY_G1	0 5 10 15 20 25	1.45	9.2e-02	6.4e-01
	REACTOME_APC_C_MEDIATED_DEGRADATION_OF_CELL_CYCLE_PROTEINS	0 5 10 15 20 25	1.45	9.2e-02	6.4e-01
	REACTOME_DISEASES_OF_GLYCOSYLATION	0 5 10 15 20 25	-1.73	5.1e-03	6.4e-01
	REACTOME_DISEASES_ASSOCIATED_WITH_GLYCOSAMINOGLYCAN_METABOLISM	0 5 10 15 20 25	-1.54	1.2e-02	6.4e-01
	DEREGULATED_CDK5_TRIGGERS_MULTIPLE_NEURODEGENERATIVE_PATHWAYS_IN_ALZHEIMER_S_DISEASE_MODELS	0 5 10 15 20 25	-1.39	9.5e-02	6.4e-01
	REACTOME_DEATH_RECEPTOR_SIGNALLING	0 5 10 15 20 25	-1.60	1.0e-02	6.4e-01
	REACTOME_CLASS_B_2_SECRETIN_FAMILY_RECEPTORS	0 5 10 15 20 25	-1.48	2.4e-02	6.4e-01



D2 sCs : Top enriched Pathways (GSEA), Old vs Young

Pathway	Gene ranks	NES	pval	padj
REACTOME_MUSCLE_CONTRACTION		2.62	3.1e-08	1.5e-06
REACTOME_TRANSPORT_OF_SMALL_MOLECULES		2.29	3.2e-05	4.7e-04
REACTOME_CARDIAC_CONDUCTION		2.31	4.8e-05	6.4e-04
REACTOME_ION_HOMEOSTASIS		2.01	4.9e-04	4.0e-03
REACTOME_LEISHMANIA_INFECTION		2.02	9.6e-04	6.9e-03
REACTOME_M_PHASE		-3.92	1.0e-10	1.2e-08
REACTOME_DNA_REPAIR		-3.88	1.0e-10	1.2e-08
REACTOME_CELL_CYCLE_MITOTIC		-4.57	1.0e-10	1.2e-08
REACTOME_CELL_CYCLE_CHECKPOINTS		-4.16	1.0e-10	1.2e-08
REACTOME_CELL_CYCLE		-4.79	1.0e-10	1.2e-08

