

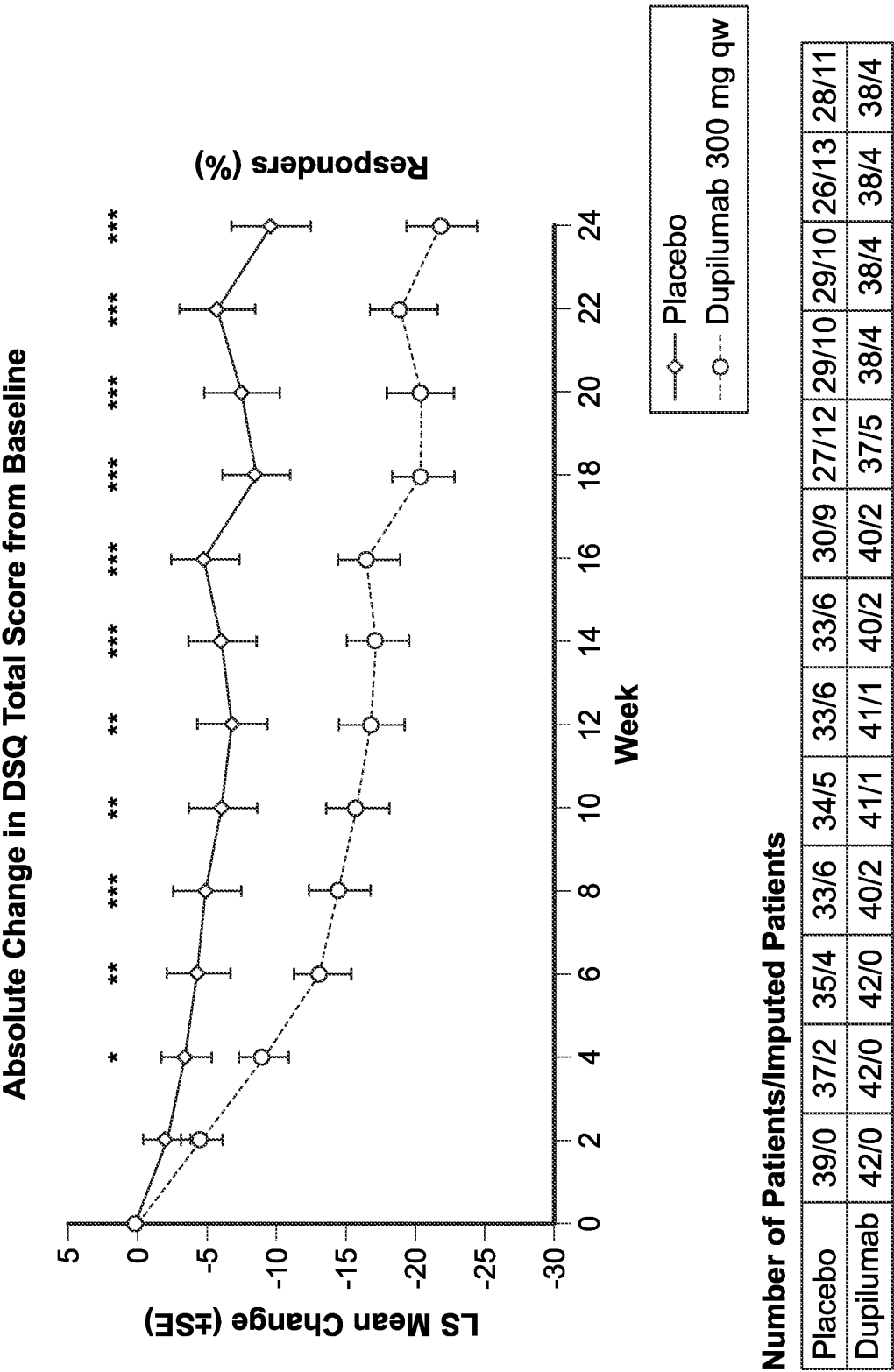


FIG.1

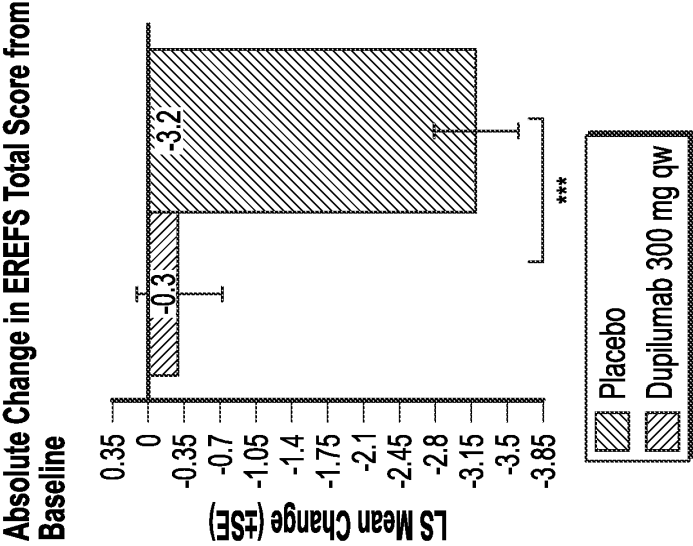
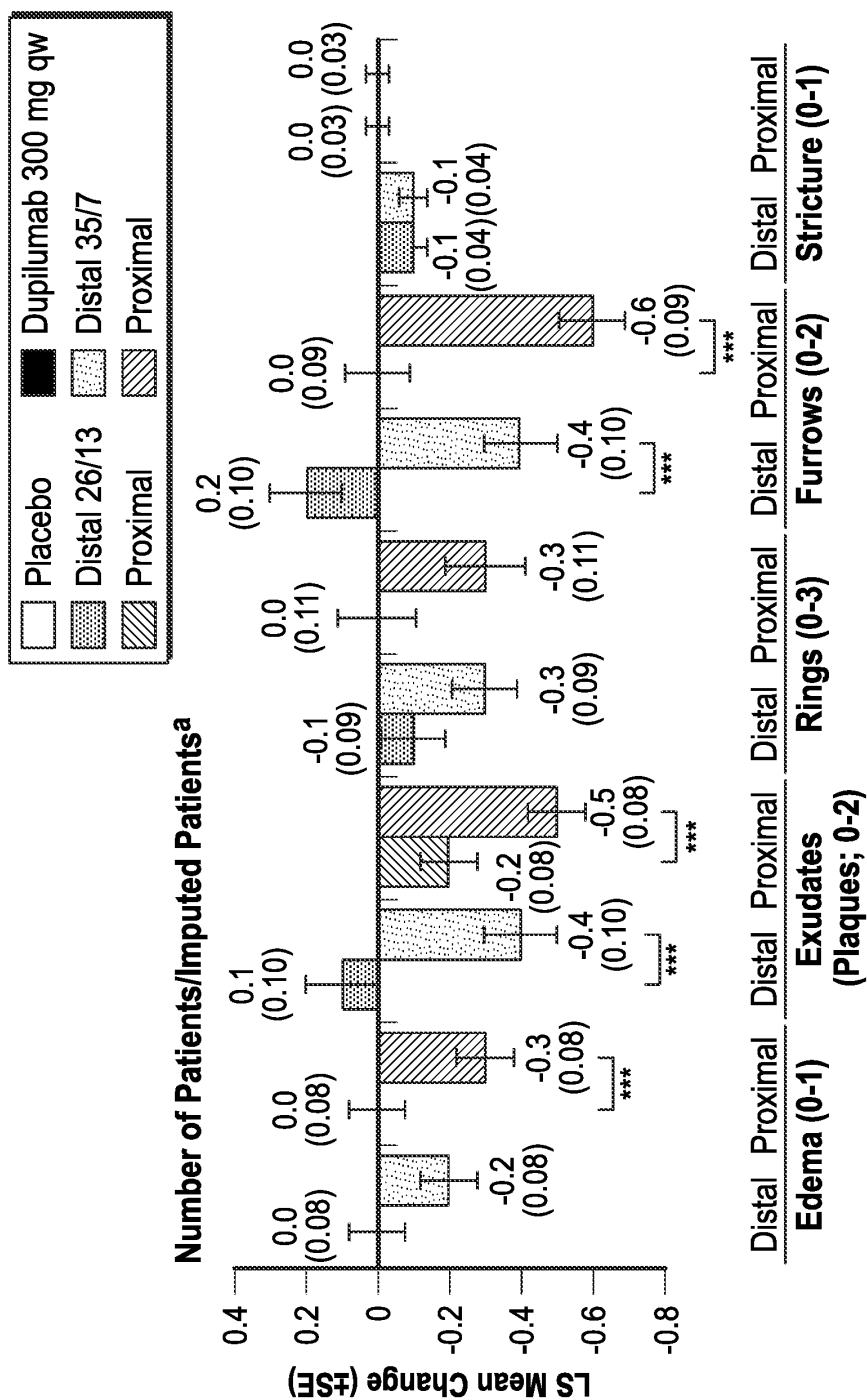


FIG. 2A



2025

Baseline mean (SD)		0.7 (0.46)	0.8 (0.40)	0.6 (0.75)	0.9 (0.80)	0.7 (0.69)	0.7 (0.81)	0.9 (0.57)	1.0 (0.64)	0.2 (0.37)	0.2 (0.38)
Distal											
Proximal		0.6 (0.49)	0.6 (0.49)	0.6 (0.68)	0.7 (0.78)	0.8 (0.77)	0.7 (0.72)	0.8 (0.51)	0.9 (0.60)	0.1 (0.34)	0.0 (0.00)

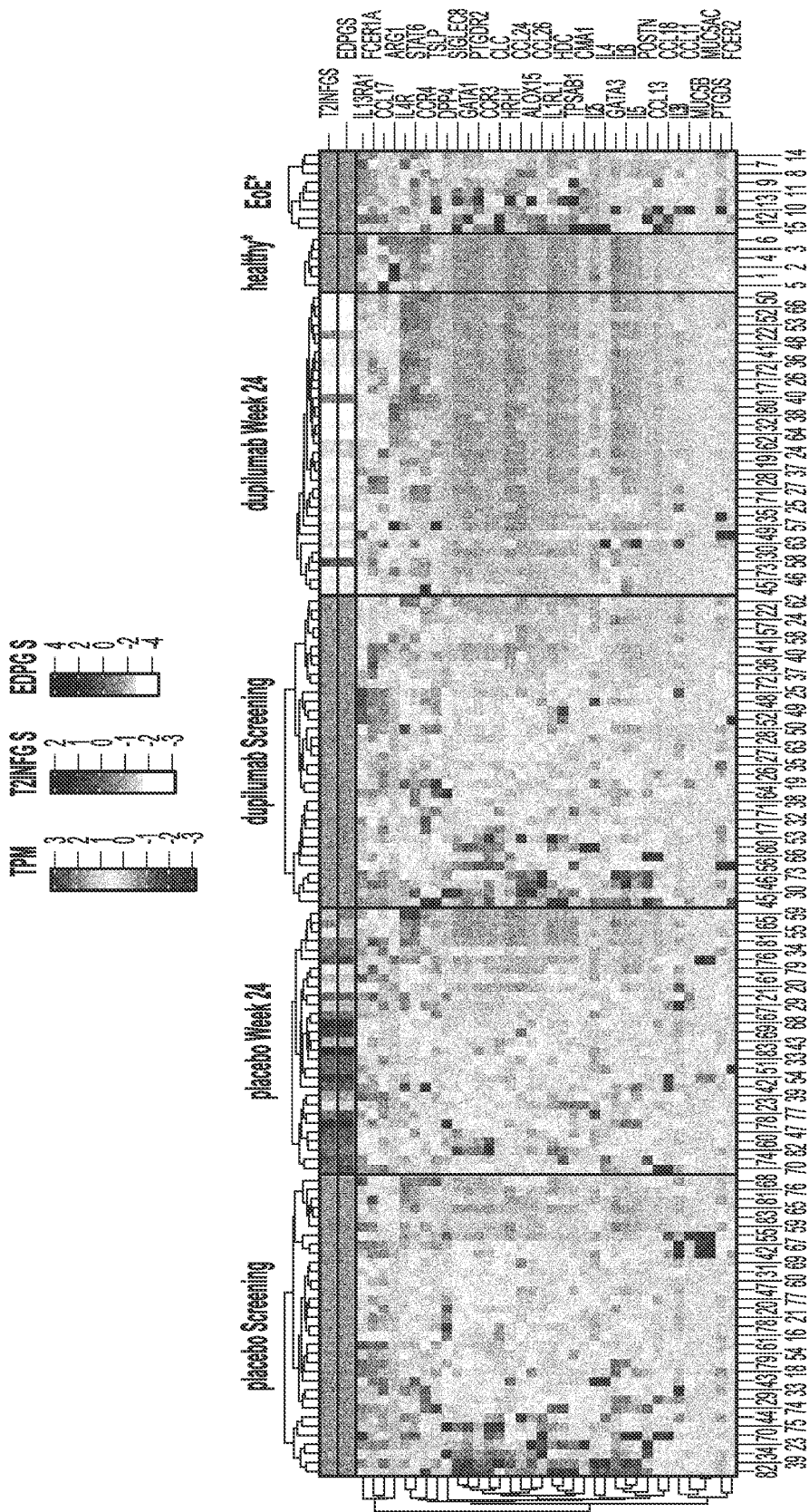


FIG. 3

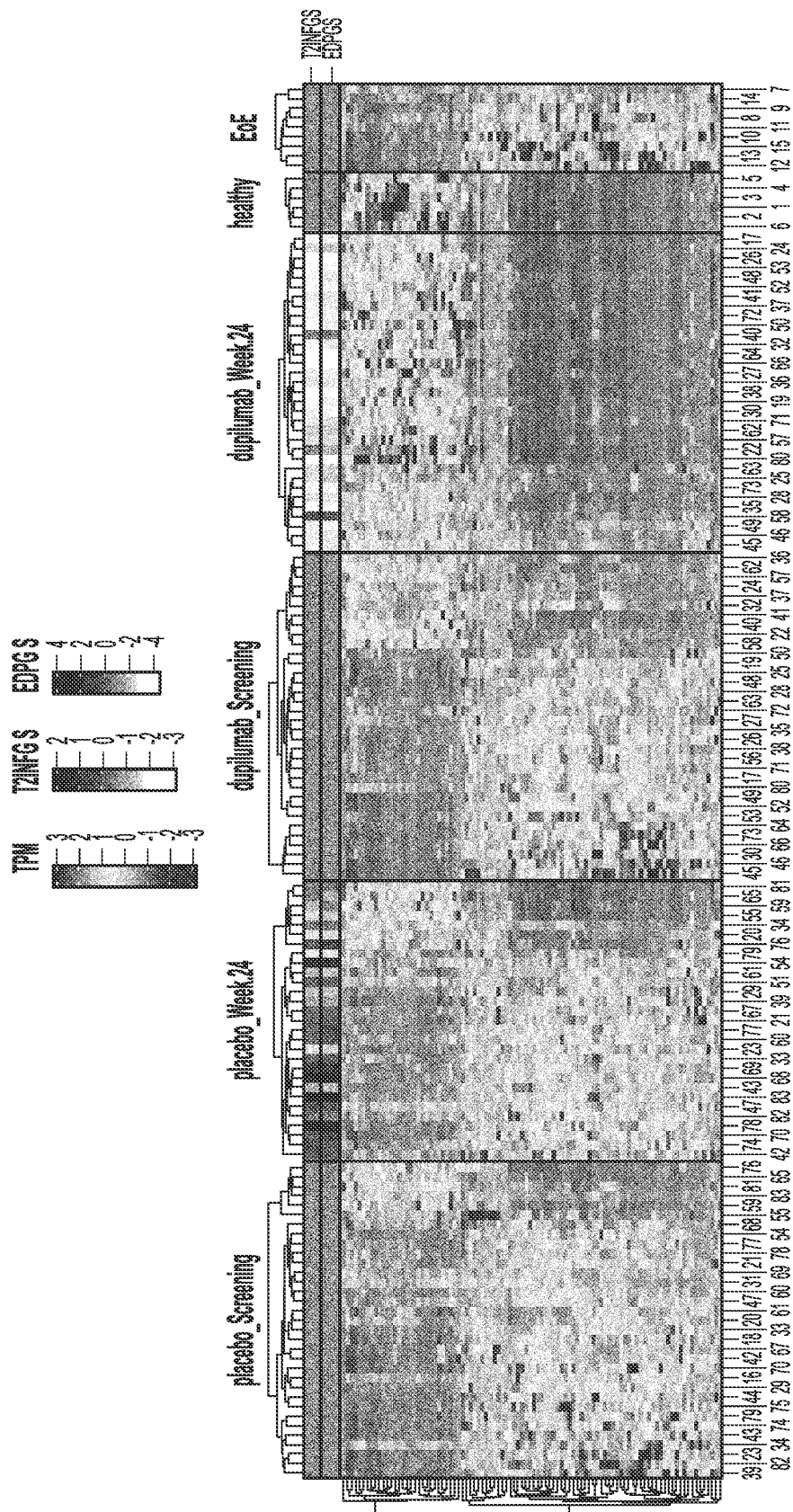


FIG. 4

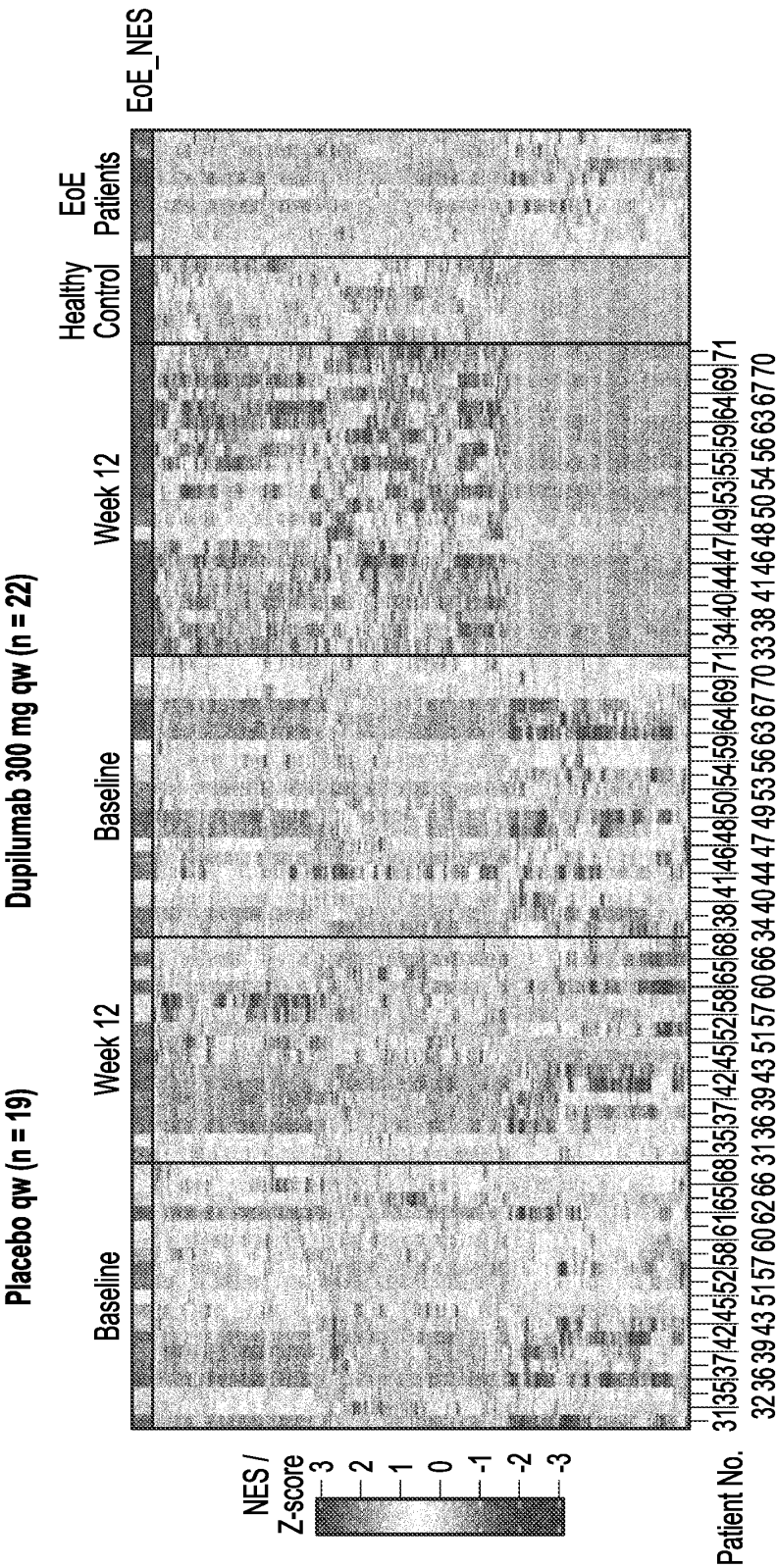


FIG. 5

Gene	EoE Transcriptome		Dupilumab 300 mg qw	
	Log ₂ Fold Change from Control	q Value	Log ₂ Fold Change from Control	q Value
TNFAIP6	9.54	3.80E-21	-9.24	3.10E-41
LRRC31	11.26	6.50E-31	-8.46	6.40E-30
SLC26A4-AS1	9.78	3.40E-46	-8.2	5.70E-29
ALOX15	9.93	8.20E-45	-8.1	3.80E-52
CCL26	8.45	4.20E-53	-6.56	6.30E-95
TGM6	7.32	4.79E-08	-6.53	2.70E-19
NRXN1	7.04	1.60E-31	-5.76	3.00E-32
PMCH	7.83	1.40E-13	-5.63	1.30E-15
SLC26A4	8.05	3.20E-33	-5.35	2.50E-27
CXCL1	6.78	2.10E-17	-5.28	5.90E-22
CCR3	6.99	2.05E-11	-5.02	4.30E-37
TREML2	5.85	2.60E-24	-4.96	1.60E-20
POSTN	8.35	4.90E-34	-4.92	6.70E-24
LURAP1L	5.72	2.30E-64	-4.67	7.10E-40
CXCL6	5.86	3.60E-12	-4.64	5.80E-20
CRTAC1	-6.76	1.10E-36	4.65	3.90E-18
BC107108	-4.11	1.00E-06	4.68	4.40E-17
SFTA2	-5.36	7.20E-07	4.99	3.10E-19
C2orf16	-1.84	4.00E-11	5.23	7.90E-17
KRTAP3-2	-7.46	1.70E-16	5.27	1.40E-19
PNLIPRP3	-5.94	1.70E-16	5.27	2.20E-15
CIDEA	-5.38	5.40E-05	5.29	4.80E-19
SLC8A1-AS1	-5.97	4.20E-20	5.41	3.20E-20
SPINK8	-6.54	4.90E-13	5.42	6.80E-19
DPCR1	-5.37	6.60E-30	5.6	2.50E-15
MUC22	-5.27	9.00E-05	5.72	1.40E-18
CRISP2	-4.75	6.50E-08	5.76	2.60E-17
DSG1	-4.09	1.70E-09	5.81	9.00E-41
GYS2	-5.03	1.10E-06	6.67	8.70E-26
CRISP3	-6.6	4.80E-10	7.63	1.10E-37

FIG. 6

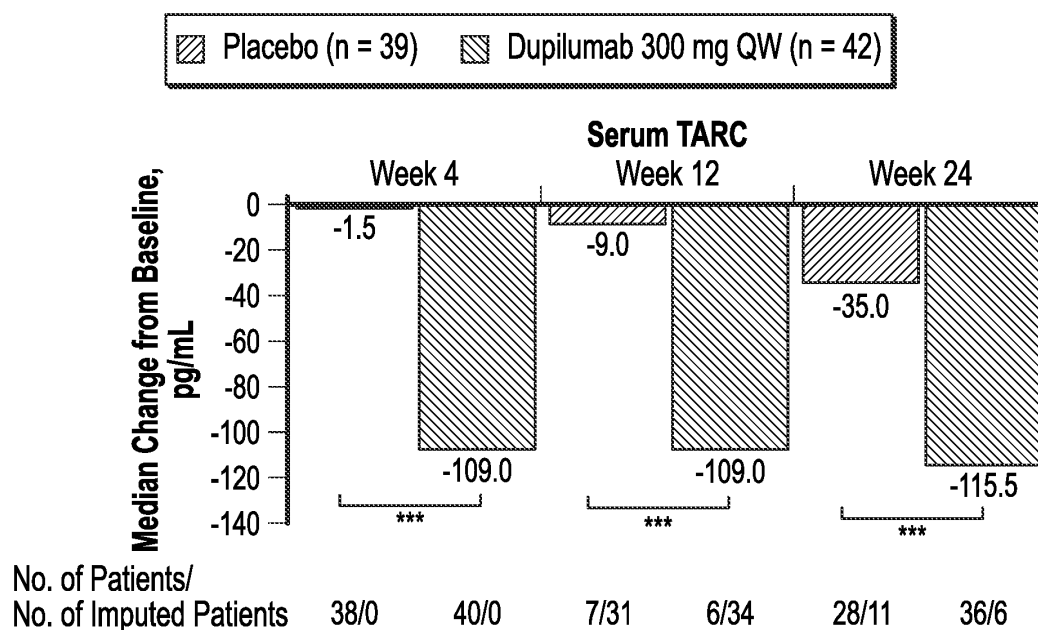


FIG. 7A

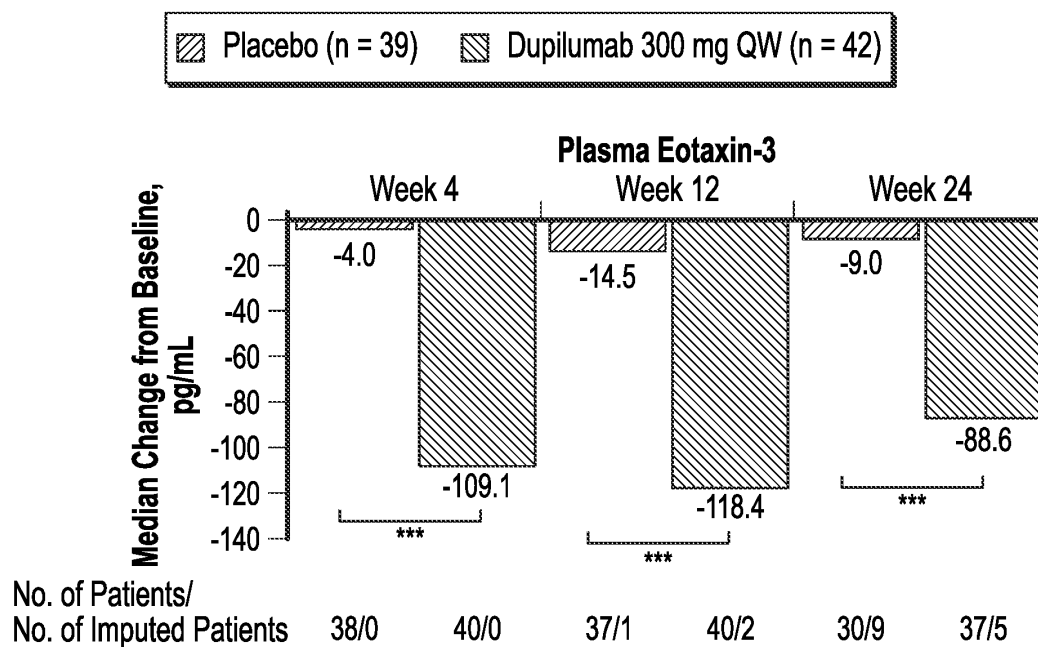


FIG. 7B

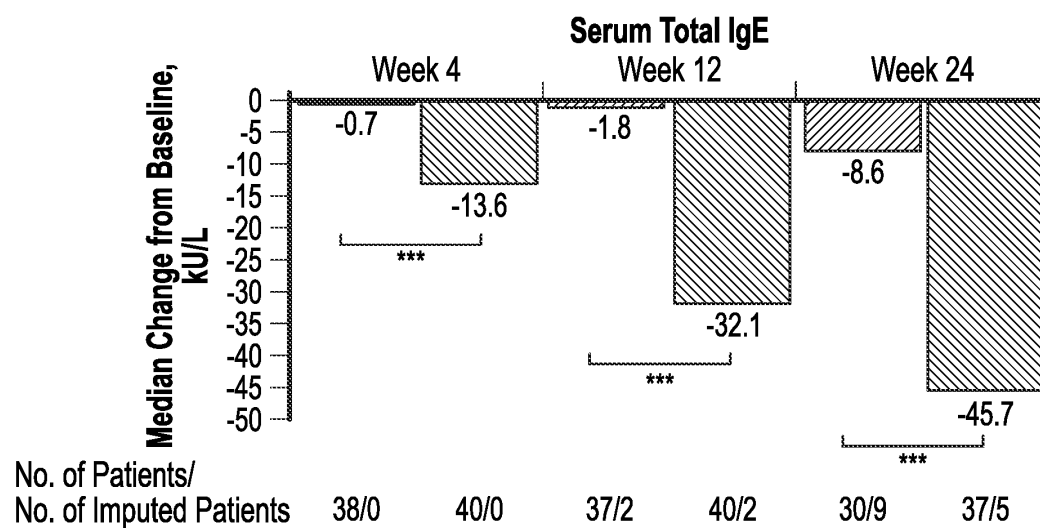


FIG. 7C