	fit_score_time $p = 1.71e-19$								
brf_gmo -	-	***	**	NS	***	***	***	***	
brf_gmonrlmf - *	***	-	***	***	***	NS	***	NS	
brf_gmosa -	**	***	-	NS	***	***	***	***	
brf_gmosanrlmf - N	NS	***	NS	-	***	***	***	***	
brf_gso - *	***	***	***	***	-	***	***	***	
brf_gsonrlmf - *	***	NS	***	***	***	-	***	***	
brf_lmo - *	***	***	***	***	***	***	-	***	
brf_lmonrlmf - *	***	NS	***	***	***	***	***	-	
-	- brt_gmo	brf_gmo_nrlmf -	brf_gmosa -	brf_gmosanrlmf -	brf_gso -	brf_gso_nrlmf -	- brf_lmo	brf_lmonrlmf -	