

Supplemental Table 16. Age-related genes in common among four or more tissues

CG_ID	No. of Tissues	Tissue ID (up- or down-regulated with age):				
CG5453	5	Acg (up) ^a	Brain (up)	MT (down) ^b	MU (up)	Tet (up)
CG14005	4	Acg (up)	Gut (down)	MT (up)	Tet (down)	
CG7220	4	Brain (up)	Fat (down)	Gut (up)	MT (up)	
CG14216	4	Brain (down)	Fat (down)	Gut (up)	MT (down)	
CG14179	4	Fat (up)	Gut (up)	MT (up)	MU (up)	
CG18445	4	Acg (up)	Gut (up)	MT (up)	MU (down)	
CG6413	4	Fat (down)	MT (up)	MU (up)	Tet (up)	
CG15797	4	Acg (up)	Gut (up)	MU (down)	Tet (down)	
CG8056	4	Acg (up)	MT (up)	MU (down)	Tet (up)	
CG18316	4	Acg (up)	Brain (down)	MT (down)	Tet (down)	
CG14150	4	Acg (up)	Brain (up)	MU (up)	Tet (up)	
CG3101	4	Acg (up)	Brain (up)	MT (down)	MU (down)	
CG7338	4	Brain (down)	Gut (up)	MU (up)	Tet (up)	
CG15009	4	Acg (up)	Fat (down)	Gut (down)	Tet (up)	
CG4237	4	Acg (down)	Brain (down)	Gut (down)	MT (down)	
CG8895	4	Acg (up)	Fat (down)	Gut (down)	Tet (up)	

^a(up) represents that the age-related gene was upregulated with age in the corresponding tissue.

^b(down) represents that the age-related gene was downregulated with age.