

## Step 1:

Check that all specimens belong to the correct sample

(if sample name is simply wrong, that will be fixed in step 2)

	specimens		samples	er_formation_name	er_specimen_alternatives	specimen_igsn	specimen_size
1	ag1-6a	belongs to	is033e				
2	ag1-6b	belongs to	is033e				
3	is031c2	belongs to	is033e				
4	is031d2	belongs to	is031d				
5	is031e2	belongs to	is031e				
6	is031f1	belongs to	is031f				
7	is032c1	belongs to	is032c				
8	is032d2	belongs to	is032d				
9	is032f1	belongs to	is032f				
10	is032i2	belongs to	is032i				
11	is033b2	belongs to	is033b				
12	is033c2	belongs to	is033c				
13	is033e2	belongs to	is033e				
14	is033g1	belongs to	is033g				
15	sc12b1	belongs to	sc12b				