Type, chosen from dropdown menu (usually "Sample")	*Autosampler location*	*Description*	*Must match filename of method that is loaded. Include everything but ".m".*	*The data file name. Must be unique or you will overwrite data.*
Type	**Vial**	**Sample**	**Method/keyword**	**Data file**
Sample	2	a-Pinene	chem472-18_method	group1_pinene001
Sample	2	a-Pinene replicate	chem472-18_method	group1_pinene002
Sample	5	Thujone	chem472-18_method	group1_thujone001
Sample	6	Limonene	chem472-18_method	group1_limonene001