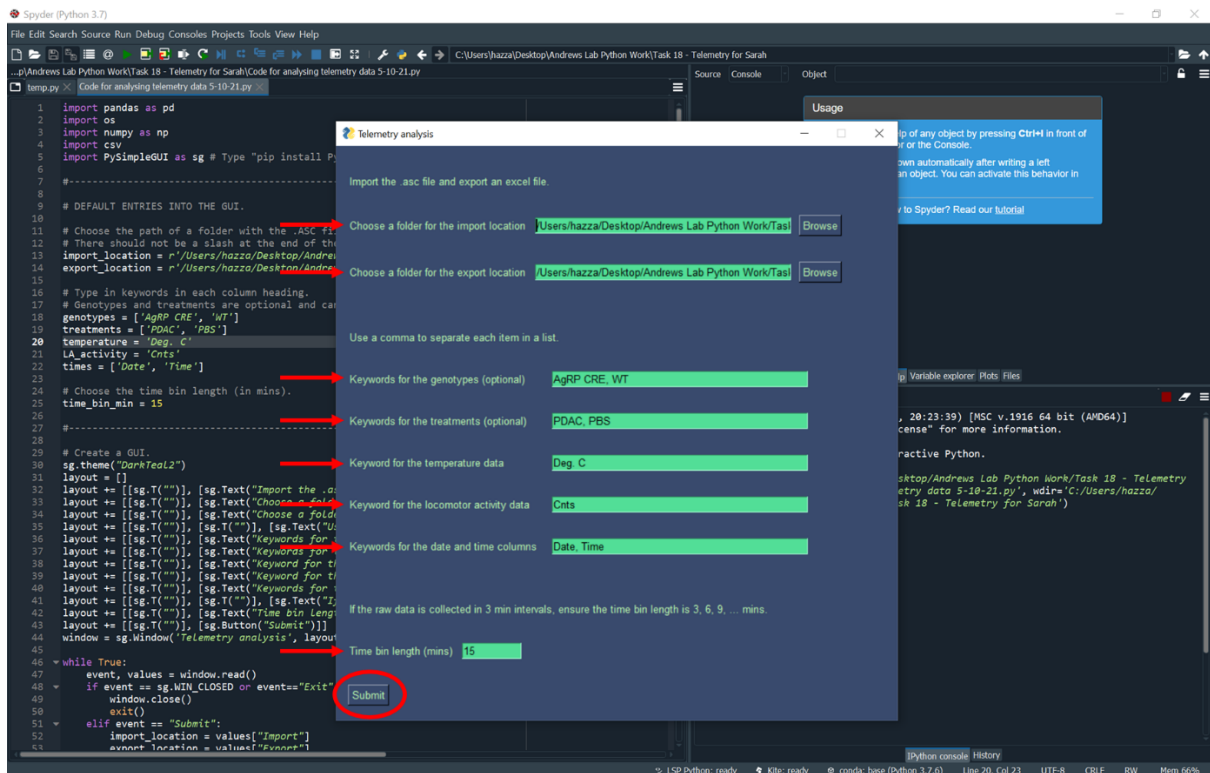


Guide to the telemetry code

Purpose: the .ASC telemetry output files have column titles that describe whether temperature or locomotor activity data is used, the mouse number and sometimes the genotype and treatment. This code separates the data based on these categories and also creates time bins.

1. Select all the options for the time bins analysis and click “submit”. Here is an explanation of all the options:



- **Import location:** the import location is a folder that contains the raw .ASC data. The code will analyse each .ASC file in the folder.
- **Export location:** the export location is a folder for the analysed files.

The column titles from the raw .ASC files need to be separated into unique keywords. See the screenshot below. Make sure different entries are also separated by a comma.

- The mouse number will be automatically sliced out from the column titles by looking for numbers.
- **Genotypes** and **treatments:** these could be 'AgRP CRE, WT' and 'PDAC, PBS' respectively. These keywords are also optional.
- **Temperature data** and **locomotor activity data:** these are usually 'Deg. C' and 'Cnts'.
- **Date** and **time:** these are usually 'Date' and 'Time'.
- **Time bin length:** this should be a multiple of the number of minutes that the raw data is collected. In the time column in the screenshot below, this data is collected in 3-minute intervals. Thus, the time bins should be 3, 6, 9, ... minutes.

AutoSave Telemetry file Harry Dempsey

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Clipboard Font Alignment Number Styles Cells Editing Analysis Sensitivity

B215 2326 AgRP CRE/PBS Deg. C Time

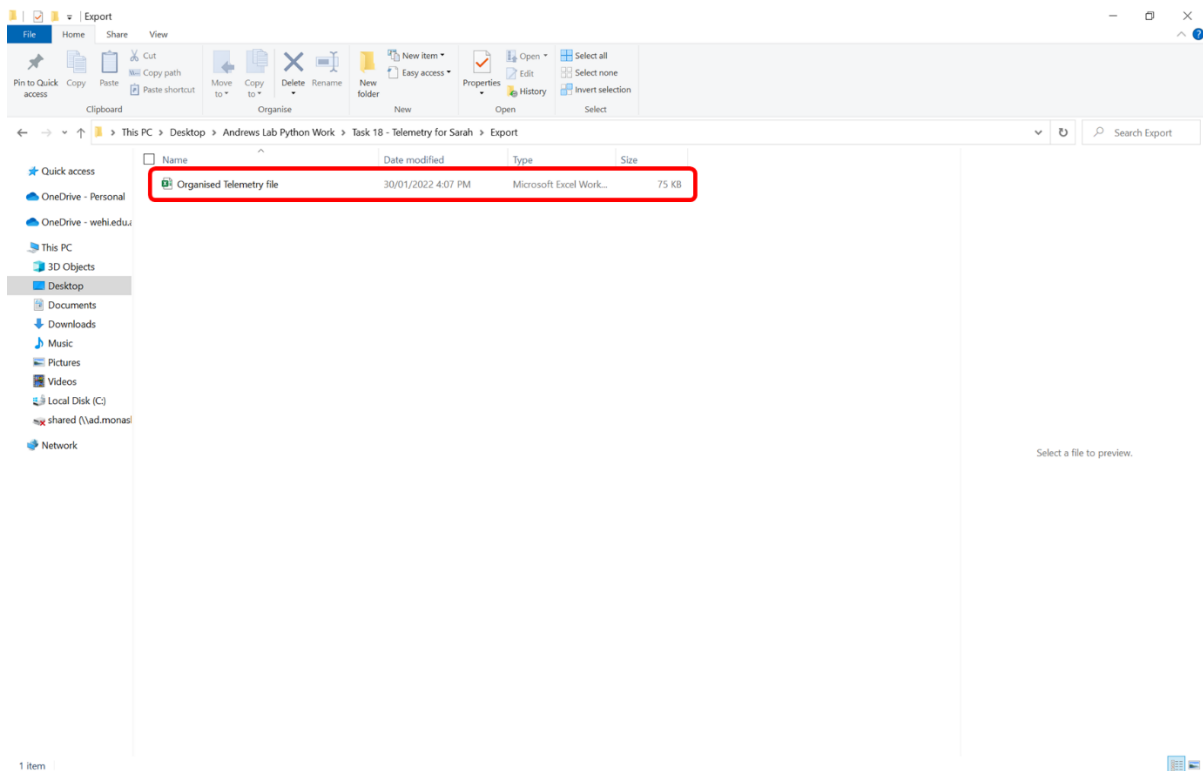
	A	B	C	D
204	High Clipping Limit: 42.000000			
205	Number Of Records: 2385			
206				
207	Animal ID: 2351			
208	Group ID: AgRP CRE/PDAC			
209	Units Measured: Cnts			
210	Sampling Interval: 00:03:00			
211	Low Clipping Limit: 0.000000			
212	High Clipping Limit: 5000.000000			
213	Number Of Records: 2385			
214				
215	2326 AgRP CRE/PBS Deg. C Date	2326 AgRP CRE/PBS Deg. C Time	2326 AgRP CRE/PBS Deg. C Data	2326 AgRP CRE/PBS C
216	05/20/20	14:07:21	NaN	
217	05/20/20	14:10:21	36.7	
218	05/20/20	14:13:21	36.53	
219	05/20/20	14:16:21	36.44	
220	05/20/20	14:19:21	36.42	
221	05/20/20	14:22:21	36.54	

Ready Accessibility: Good to go 190%

Annotations:

- Mouse number (points to column B header)
- Genotype (points to column C header)
- Treatment (points to column D header)
- Temperature or locomotor activity data (points to column E header)
- Date, time or data column (points to column F header)

2. Go to the export location to find the time binned file.



3. Here is the exported data.

- The locomotor activity data and temperature data are split into 2 sheets.
- The column headings within each sheet are labelled with mouse number, genotype and treatment (if listed). They are arranged by genotype first and then treatment.
- The time bins in minutes since the first time point and the corresponding dates and times are included. Assuming that the dark cycle is between 7pm and 7am, these dates and times are highlighted in grey.

Locomotor activity sheet

Organised Telemetry file									
Harry Dempsey									
File Home Insert Draw Page Layout Formulas Data Review View Help									
Clipboard Font Alignment Number Styles Cells Editing Analysis Sensitivity									
A1 X fx Date and time									
	A	B	C	D	E	F	G	H	I
1	Date and time	Time bins (mins)	2326 AgRP CRE/PBS Cnts Data	2327 AgRP CRE/PBS Cnts Data	2329 AgRP CRE/PBS Cnts Data	2333 AgRP CRE/PBS Cnts Data	2336 AgRP CRE/PDAC Cnts Data	2341 AgRP CRE/PDAC Cnts Data	2349 AgRP CRE/PDAC Cnts Data
2			2326	2327	2329	2333	2336	2341	2349
3			AgRP CRE	AgRP CRE	AgRP CRE	AgRP CRE	AgRP CRE	AgRP CRE	AgRP CRE
4			PBS	PBS	PBS	PBS	PDAC	PDAC	PDAC
5	2020-05-20 14:07:21	0	0	0	0	0	0	0	0
6	2020-05-20 14:22:21	15	495	516	315	546	462	560	560
7	2020-05-20 14:37:21	30	122	100	18	445	0	757	757
8	2020-05-20 14:52:21	45	42	36	22	139	88	215	215
9	2020-05-20 15:07:21	60	16	46	46	33	35	30	30
10	2020-05-20 15:22:21	75	36	11	79	26	2	86	86
11	2020-05-20 15:37:21	90	65	531	21	54	283	58	58
12	2020-05-20 15:52:21	105	11	122	12	8	15	35	35
13	2020-05-20 16:07:21	120	114	94	4	100	36	11	11
14	2020-05-20 16:22:21	135	262	50	23	344	9	58	58
15	2020-05-20 16:37:21	150	73	188	273	364	167	218	218
16	2020-05-20 16:52:21	165	32	76	12	12	181	163	163
17	2020-05-20 17:07:21	180	0	486	35	23	415	75	75
18	2020-05-20 17:22:21	195	86	398	305	21	30	9	9
19	2020-05-20 17:37:21	210	46	63	14	30	0	33	33
20	2020-05-20 17:52:21	225	264	82	18	51	0	387	387
21	2020-05-20 18:07:21	240	494	100	20	235	9	261	261
22	2020-05-20 18:22:21	255	459	426	170	196	147	0	0
23	2020-05-20 18:37:21	270	105	157	330	159	576	33	33
24	2020-05-20 18:52:21	285	20	25	29	55	163	57	57
25	2020-05-20 19:07:21	300	4	140	54	95	308	190	190
26	2020-05-20 19:22:21	315	198	329	277	336	468	97	97
27	2020-05-20 19:37:21	330	206	395	230	162	636	600	600
28	2020-05-20 19:52:21	345	413	385	431	393	814	545	545
29	2020-05-20 20:07:21	360	426	319	244	471	969	441	441
30	2020-05-20 20:22:21	375	133	589	439	414	584	501	501
31	2020-05-20 20:37:21	390	2	567	389	230	773	383	383
32	2020-05-20 20:52:21	405	83	652	451	578	719	538	538
33	2020-05-20 21:07:21	420	36	710	186	300	708	823	823
34	2020-05-20 21:22:21	435	38	403	47	188	717	488	488
	Locomotor activity data	Temperature data							

Temperature data sheet

Date and time	Time bins (mins)	2326 AgRP CRE/PBS Deg. C Data	2327 AgRP CRE/PBS Deg. C Data	2329 AgRP CRE/PBS Deg. C Data	2333 AgRP CRE/PBS Deg. C Data	2336 AgRP CRE/PD Deg. C Data	2341 AgRP CRE/PD Deg. C Data
2020-05-20 14:07:21	0	0	0	0	0	0	0
2020-05-20 14:22:21	15	36.526	36.114	37.556	36.798	36.032	36.4
2020-05-20 14:37:21	30	36.29	35.64	36.802	36.83	35.022	36.4
2020-05-20 14:52:21	45	35.748	34.858	36.772	36.328	35.332	36
2020-05-20 15:07:21	60	35.546	34.644	36.722	35.49	35.468	35.1
2020-05-20 15:22:21	75	35.51	34.794	36.676	34.782	35.406	35
2020-05-20 15:37:21	90	35.592	35.16	36.756	34.962	35.49	34.4
2020-05-20 15:52:21	105	35.58	35.198	36.574	34.65	35.006	34.4
2020-05-20 16:07:21	120	35.648	34.8	36.672	34.874	35.488	35.1
2020-05-20 16:22:21	135	36.152	34.93	36.38	36.004	35.14	34.4
2020-05-20 16:37:21	150	35.902	34.876	36.748	36.46	35.25	35.4
2020-05-20 16:52:21	165	35.386	34.95	36.698	36.522	35.34	35.1
2020-05-20 17:07:21	180	35.79	35.326	36.968	35.87	35.812	35.1
2020-05-20 17:22:21	195	35.942	35.65	36.978	35.31	35.288	35
2020-05-20 17:37:21	210	35.72	35.414	36.584	34.892	34.838	34.4
2020-05-20 17:52:21	225	36.262	34.96	36.752	34.762	34.918	35.4
2020-05-20 18:07:21	240	36.428	34.924	36.748	35.294	35.108	35.1
2020-05-20 18:22:21	255	36.706	35.304	36.982	36.104	35.218	35.1
2020-05-20 18:37:21	270	36.752	35.678	37.634	36.714	36.37	35.1
2020-05-20 18:52:21	285	35.972	35.398	37.35	36.556	36	34.4
2020-05-20 19:07:21	300	35.63	35.552	37.204	36.828	35.744	35.1
2020-05-20 19:22:21	315	35.926	36.16	37.52	36.796	36.598	35.4
2020-05-20 19:37:21	330	36.368	36.624	38.132	36.572	36.808	36.1
2020-05-20 19:52:21	345	36.844	36.786	38.478	36.656	37.36	37.4
2020-05-20 20:07:21	360	37.102	36.912	38.506	36.86	37.412	37
2020-05-20 20:22:21	375	37.148	37	38.652	37.102	37.432	37.1
2020-05-20 20:37:21	390	36.456	37.028	38.584	37.21	37.43	37.1
2020-05-20 20:52:21	405	36.08	37.026	38.628	37.362	37.374	37.1
2020-05-20 21:07:21	420	36.172	37.178	38.58	37.37	37.434	37.1
2020-05-20 21:22:21	435	35.982	36.968	38.214	36.874	37.352	37.1

- You can also change the default values that appear in the GUI by modifying this section of the code. This is in the “Create_GUI.py” file and under the function “default_values”. This is useful if you need to run the code multiple times with similar options. Make sure that the input formats are kept the same. For example:

- Folder paths should have forward slashes, no slash at the end and be contained in ‘ ’ marks.
- genotypes, treatments and times should have this format: [‘input1’, ‘input2’]
- temperature and LA_activity should have this format: ‘input’

Spyder (Python 3.7)

```
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C:\Users\hazza\Desktop\Github final\Codes\Telemetry_analysis\Codes\Create_GUI.py

Create_GUI.py* x Create_time_bins.py x Run_program.py x

1 import PySimpleGUI as sg
2
3 def default_values():
4
5     default = {}
6
7     # Choose the path of a folder with the .ASC file, so the code can import every
8     # file in the folder.
9     # There should not be a slash at the end of these folder paths.
10    default['Import Location'] = r'/Users/hazza/Desktop/Telemetry code/Import'
11    default['Export Location'] = r'/Users/hazza/Desktop/Telemetry code/Export'
12
13    # Type in keywords in each column heading.
14    # Genotypes and treatments are optional and can be left as [].
15    default['Genotypes'] = ['AgRP CRE', 'NT']
16    default['Treatments'] = ['PDAC', 'PBS']
17    default['Temperature'] = 'Deg. C'
18    default['LA_activity'] = 'Cnts'
19    default['Times'] = ['Date', 'Time']
20
21    # Choose the time bin length (in mins).
22    # If the raw data is collected in 3 min intervals, ensure the time bin length is
23    # 3, 6, 9, ... mins.
24    default['Time bin (mins)'] = 15
25
26    return(default)
27
```