

kConFab cost recovery schedule

Jan 2017

In the past, kConFab have charged approved projects an Administration Fee (in part because the NHMRC insist that we recover some costs) but this was replaced in 2011 in favour of a more transparent, unit-based system. The kConFab Executive agreed that every year the default increase will be in line with the Australian CPI for all biospecimens and data, to be confirmed annually by the Executive Committee.

These updated charges are shown in the table below. For this financial year (July 1 2016 – June 30, 2017) billing will be in early July.

The biospecimen charges are roughly based on what other Australian biobanks charge and what we think people can afford to pay, rather than on what it costs to generate them (else a lot of projects would be far too expensive for any researcher to pay for). Our estimation is that with this scheme some researchers will end up paying more, and some will pay less, and the overall income for kConFab will be slightly, but not hugely increased. Some items such as LCLs, will have a sliding scale to avoid some projects being priced out of existence, and LCLs benefit more than just the project for which they are generated.

The data charges are a tiny fraction of what it costs to obtain all the composite data, and are not a direct reflection of the time it takes to fulfil a data request. We have grouped the data into blocks of: Base data (basic person/family data including DOB, DOD and cancer details), Pathology data, Mutation details, Epidemiology data, Treatment/prophylactic surgery data, Medicare/PBS, Pedigree data and Contact details (which are only given to the CFU and psychosocial projects). The data items in each of these blocks are listed below. Please note that:

- Base data would only be charged for the data-only projects.
- If you request blood, cell lines or DNA then you don't get charged for base or mutation data; if you request tissue you don't get charged for base or pathology data.
- One hour of data processing time by the Data Manager is included with each data group requested. Any time spent above that will be charged extra (\$250/hour) to cover the extra work of a subset of requests. This charge is a reflection of the large amount of data, from many sources, being given out for these requests, and will help contribute to the costs of collecting all the data from each individual (estimated to be about \$750 in total, per person).
- If data updates are requested under an approved project, the researcher will be charged an hourly rate of \$250/hour with a minimum of 1 hour being billed. It is suggested that a quote for the estimated time to be taken should be asked for before the data manager commences this work.

These prices below are those that we will charge for projects on which the Principal Investigator is based in Australia. For projects led by an overseas investigator, but which are clearly genuine collaborations with an investigator in Australia, we will charge 1.5X these rates, and for applications from Principal Investigators overseas which do not include any researchers in Australia, we will charge 2X these rates. Companies will be charged at least 2.5X times these rates (subject to negotiation).

We should also forewarn you that there might be further changes next year, depending on our financial situation and the demands of funding agencies, but in the meantime we hope that these charges will help you to write grant applications to use kConFab data and biospecimens, and prepare you for a change in invoicing next year.

Lastly, there will be no charges for data queries necessary to write a grant application (unless they involve an extraordinary amount of Eveline's time e.g. more than two days work).

Georgia Chenevix-Trench (Chair, kConFab Finance sub-committee)

Unit Cost in Australian dollars	Unit
3.1	≤1ug DNA
6.1	>1 and ≤3ug DNA
12.2	>3 and ≤5ug DNA
24.3	>5ug DNA
by negotiation	>10ug DNA
14.2	Plasma (n=1-5)
11.1	Plasma (n=6-50)
9.1	Plasma (n>50)
164.1	Cell Line (n=1-5)
82.1	Cell lines (n=6-50)
35.5	Cell lines (n>50)
20.3	Viable White Blood Cells frozen in 10% DMSO/FCS (n=1-5)
15.2	Viable White Blood Cells frozen in 10% DMSO/FCS (n=6-50)
10.1	Viable White Blood Cells frozen in 10% DMSO/FCS (n=50+)
121.6	TMA Section
152	FFPE Block to borrow/core
10.1	FFPE stained slide to keep (3-5 microns)
4.1	FFPE slide to borrow or view scanned image
12.2	FFPE unstained slide (3-5 microns)
24.4	FFPE unstained slide (8-10 microns) <i>price to be discussed for >10 microns</i>
704	Whole fresh tissue, including shipping
354.6	Whole fresh tissue, with no shipping costs
212.7	Fresh frozen tissue (1-10 sections)
106.4	RNA Later (1-10 sections)
12.2	Shavings from fresh frozen tissue
12.2	Slide - stained (fresh frozen) (3-5 microns)
12.2	Slide - unstained (fresh frozen) (3-5 microns)
1595.5	CASCADE rapid autopsy tissue (fresh/frozen)
2.1	Base Data for first 10,000 consented or deceased UPNs
1.5	Base Data for next 10,000 consented or deceased UPNs
1.1	Base Data for next 10,000 consented or deceased UPNs
1.1	Base Data for first 30,000 non-consented UPNs
0.76	Base Data for next 30,000 non-consented UPNs
0.51	Base Data for next 30,000 non-consented UPNs
2.1	Consent form per UPN
2.1	Pathology Data per UPN
2.1	Mutation Data per UPN
2.1	Epi Data per UPN
6.1	Treatment Data per UPN
2.1	Family Relationships per UPN (except parent ids)
2.1	Contact information per UPN
0.51	SNP data per UPN for 1-5 SNPs
1.51	SNP data per UPN for 6-99 SNPs
3.6	SNP data per UPN for 100-210,000 SNPs
12.2	SNP data per UPN for >210,000 SNPs
253.3	Per extra hours beyond the 1 hour usually needed to fulfil a data request

For a complete list of available data fields see <http://www.kconfab.org/Collection/DataFields.shtml>

Table	Data Category
Address	Contact
Blood Specimen	Base
Cancer	Base
Death	Base
Family	Base
Person	Base
Family Relationship (parent ids only)	Base
Family Relationship	Family Relationship
Tissue Collection	Base
Epidemiological Questionnaire	Epi
Family Mutation	Mutation
Mutation Testing	Mutation
Pathology	Pathology
QA Data for fresh tissue	Pathology
Slide Review	Pathology
Treatment	Treatment