# costseq

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# 1 Full economic cost [0/4]

1.1 [] Cost sources

#### 1.1.1 Data generation

- 1. labor
- 2. management
- 3. utilities
- 4. sequencing machines and other equipment
- 5. low level squence informatics (e.g. machine level base calls)
- 6. data submission and storage
- 7. indirect costs

### 1.1.2 Data analysis

- 1. quality assessment
- 2. technology development
- 3. downstream bioinformatics analysis
- 4. data storage and analysis

## 1.2 [] Cost information from YCGA (CMG Grant)

- 1.2.1 Dell PowerEdge R720 with 512GB of memory and 3.6TB of disk space for programming and data processing.
  - 1. \$10,000 (one time purchase in Year 1)

#### 1.2.2 New technology development costs

- 1. \$44,800
  - (a) \$11,200 per year for:
    - i. PCR plates, pipette tips, 15 ml and 50 ml Falcon tubes, 1.5ml and 2ml Eppendorf tubes, 5-20ml serological pipettes, Pasteur pipettes, patient sample collection tubes, gloves, reagent reservoirs and other disposables.

- 2. \$115,688
  - (a) \$28,922 per year for:
    - i. agarose, tris, phenol, chloroform, molecular biology grade ethanol, proteinase K, PCR buffers, Taq polymerase various chemicals and R&D sequencing lanes.
- 3. Our total R&D budget constitutes  $^{5\%}$  of the total cost of the grant, which we believe is conservative.

#### 1.2.3 In vitro functional studies

- 1. \$400,000
  - (a) 100,000/year for years 1-4

# 1.2.4 Acquisition of sample collections from other centers and NIH institutes.

1. 20,000 (over all 4 years)

#### 1.2.5 Exome capture and sequencing

- 1. \$7,656,000 in direct costs is requested over the four year period to capture and sequence over 26,000 exomes.
  - (a) Year one: At the rate of \$330 in direct costs/exome, we request \$1,914,000 in direct costs towards capture and sequence analysis of 5,800 exomes.
  - (b) Year two: At the rate of \$305 in direct costs/exome, we request \$1,914,000 in direct costs towards capture and sequencing of 6,275 exomes.
  - (c) Year three: At the rate of \$291 in direct costs/exome, we request \$1,914,000 in direct costs to capture and sequence 6,577 exomes
  - (d) Year four: At a rate of \$254 in direct costs/exome, we request \$1,914,000 in direct costs to capture and sequence 7,535 exomes.

Table 1: Exome Sequencing Costs for the duration of the funding period \*

Description	Year 1	Year 2	Year 3	Year 4
Capture and Library Preparation cost				
NimbleGen V2 Capture Oligo pools	\$36	\$34	\$32	\$32
Labor	\$57	\$54	\$51	\$51
Reagent and Supplies	\$50	\$48	\$45	\$43
Instrument Depreciation and Maintenance	\$12	\$12	\$12	\$0
Administration and IT etc. $(3\%)$	\$5	\$4	\$4	\$4
Sub Total Library Prep	\$160	\$152	\$145	\$131
Sequencing Cost				
Labor	\$6	\$5	\$5	\$5
Sequencing Reagents & Supplies	\$118	\$105	\$100	\$100
Instrument Depreciation	\$24	\$22	\$22	\$0
HPC and IT	\$8	\$6	\$5	\$3
Maintenance contract	\$10	\$10	\$10	\$10
Administrative cost $(3\%)$	\$5	\$4	\$4	\$4
Sub Total Sequencing	\$171	\$153	\$146	\$123
Total Direct cost (Library prep + Sequencing)	\$330	\$305	\$291	\$254
Indirect costs (Equipmentnot included	\$196	\$180	\$171	\$169
Direct + Indirect Cost	\$526	\$485	\$463	\$423

\*Costs are determined for 2x100 bp sequencing using Illumina HiSeq 4000

#### 1.2.6 Whole genome sequencing

- 1. \$1,404,000 in direct costs is requested over the four year period to library preparation and sequencing of 672 genomes.
  - (a) Year one: At the rate of \$2,340 in direct costs/genome, we request \$351,000 in direct costs towards library preparation and sequence analysis of 150 genomes.
  - (b) Year two: At the rate of \$2,226 in direct costs/genome, we request \$351,000 in direct costs to capture and sequence 157 genomes.
  - (c) Year three: At the rate of \$2,121 in direct costs/genome, we request \$351,000 in direct costs to capture and sequence 165 genomes.
  - (d) Year four: At a rate of \$1,754 in direct costs/genome, we ask \$351,000 in direct costs to capture and sequence 200 genomes. Table 2 : Whole-Genome Sequencing cost for the entire funding period

Description	Year 1	Year 2	Year 3	Year 4
Library Preparation Costs				
Labor	\$70	\$56	\$45	\$36
Reagent and Supplies	\$75	\$71	\$68	\$64
Instrument Depreciation	\$10	\$10	\$10	\$0
Service Contract IT	\$8	\$8	\$8	\$8
Administration $(3\%)$	\$5	\$4	\$4	\$3
Sub Total Library Prep	\$168	\$149	\$134	\$111
Sequencing Costs (1 genome/lane $2x150$ bp)				
Labor	\$82	\$78	\$74	\$70
Sequencing Reagents and Supplies	\$1767	\$1679	\$1595	\$1515
Instrument Depreciation	\$175	\$175	\$175	\$0
HPC IT	\$75	\$75	\$75	\$0
Maintenance Contract	\$10	\$10	\$10	\$10
Administrative cost $(3\%)$	\$63	\$60	\$58	\$48
Sub Total Sequencing	\$2172	\$2077	\$1987	\$1643
Total Direct Cost (Library prep + Sequencing)	\$2340	\$2226	\$2121	\$1754
Indirect Cost (Equipment cost is not included)	\$1433	\$1358	\$1287	\$1167
$\mathrm{Direct}+\mathrm{Indirect}\;\mathrm{costs}$	\$3773	\$3584	\$3408	\$2921

\*Costs are determined for 2x150 bp sequencing using Illumina HiSeq 4000  $\,$ 

# 1.2.7 Notes:

- 1. Labor costs for WGS projected to drop while exome sequencing labor costs level off.
- 2. Decrease in WGS sequencing prep is primarily from labor costs and to a lesser extent reagent costs.

### 1.2.8 Variant confirmation via Sanger Sequencing

- 1. \$60,000
  - (a) \$3.25 for the primer pair and \$1.75 for the actual Sanger sequencing. In total, we will have 12,000 variants from 1,500 exome sequencing variants.

# 1.3 [] Information from the NHGI

#### 1.3.1 Email sent to Kris Wetterstrand.

1. She said she is happy to share whatever information she has but is busy and will get back to us in a few days.

# 1.4 [] Information from Illumina

### 1.4.1 Email sent to Crane Harris.

- 1. He shared this link: http://www.molecularecologist.com/next-gen-fieldguide-2014/
- 2. "Illumina is running three regional sales meetings around the world over the next two and a half weeks, and the folks who might be able to help here are consumed with prep and travel for those. If as you get into your analysis there is a call for more system-specific information – high throughput/low throughput, clinical/research, depth of coverage, human/other, targeted/whole genome – let me know and we can look for something more targeted."