### Protocol of the exome enrichment of library pools for WES sequencing on MGI

	Title	Manufacturer	P/N
1	80% ethanol	Constanta-Pharm M" LLC	LS-002430
2	mQ water	Millipore	
3	Qubit dsDNA HS Assay Kit	ThermoFisher	IQ32854
4	Qubit dsDNA BR Assay Kit	ThermoFisher	Q32853
5	Agencourt AMPure XP	Beckman	A63881
6	Set for Target Enrichment / SSELXT Human All Exon V6, 16 rxn	Agilent Technologies	5191-4004
7	Magnetic particles with streptavidin Dynabeads MyOne Streptavidin C1 (10mg/ml)	ThermoFisher	65001
8	COT Human DNA (500µg/packaging)	Roche	11581074001
9	SALMON SPERM DNA SOLN 5 X 1ML (10 mg/ml) reagent	Thermo Invitrogen	15632011
10	RNAase inhibitor Super Rase In, 2500 units. (20U/mkl)	Thermo Invitrogen	AM2694
11	Reagent 50X DENHARDTS SOLUTION 100ML	Thermo Invitrogen	750018
12	KAPA HiFi polymerase HS	KAPA biosystems	7958897001
13	Sodium chloride and sodium phosphate 20 X SSPE 1L	Thermo Invitrogen	AM9767
14	Reagent 0.5 M EDTA PH 8.0 100ML (BUF KIT)	Thermo Invitrogen	AM9260G
15	5M NACL 500 ML EACH Reagent	Thermo Invitrogen	AM9759
16	Reagent 10% SDS SOLUTION 500 ML EACH	Thermo Invitrogen	AM9822
17	20X SSC BUFFER 500 ML EACH Reagent	Thermo Invitrogen	AM9770
18	Tris-HCl 1M 7.5 pH, 1L	Thermo Invitrogen	15567027
19	High Sensitivity DNA reagents	Agilent Technologies	G2938-85004
20	Agarose		
20	Ethidium bromide		
21	TAE buffer (x1)	Eurogen	PB022

### Reagents

Oligonucleotide s	Concentration	Sequence 5'-3
MGIAd_PCR_2	30 µM	TGTGAGCCAAGGAGTTG
MGIAd_PCR_1	30 µM	/5Phos/GAACGACATGGCTACGA
IGMblock11	1000 μM	GAACGACA+TGGC+TACGA+TCCGAC+TT
IGMblock12	1000 μM	TGTGAGCC+AAGG+AGTTGiiiiiiiiiiiiiiiiiiiiiiiiiiiii

#### Materials

	Title	Manufacturer	P/N
1	LoBind 1.5 ml microtubes	Eppendorf	22431021
2	Falcon 50 ml tubes	Eppendorf	Epp 0030 122.178
3	Filter tips 20-300	SSI	SSI-4928NSFS
4	Filter tips 2-20	SSI	SSI-4238NAFS
5	Handpieces with filters 1-10	SSI	SSI-4138NSFS*
6	Filter tips 300-1200	SSI	SSI-4348NSFS
7	Microtubes 0.5 ml	Cornig	PCR-05-C
8	Tubes 1.5ml 500 pcs/upe	SSI	SSI-1260-00
9	SSI-3111 strips and flat caps	SSI	SSI-3105*
10	Strips Eppendorf 0.2 ml	Eppendorf	Epp 0030 124.359
11	0.2 PCR tubes with ind. lids	SSI	SSI

### Equipment

	Title	Manufacturer	P/N
1	Qubit	Thermo	
2	Vortex Microspin FV-2400	Biosan	
3	Automatic pipettes Eppendorf in the range of volumes from 0.5 to 10 µl, from 2 to 20 µl, from 200 to 200 µl, from 100 to 1000 µl	Eppendorf	
4	Racks for 1.5 ml, 0.5 ml, 0.2 ml tubes	Helicon	
5	Amplifier 96-well with manually opened lid		
6	Orbital shaker, thermal shaker	Biosan	
7	Thermostat	DNA technology	
8	Bioanalyzer 2100	Agilent	
9	Magnetic rack for 1.5 ml tubes and 0.2 ml tubes	Thermo, Helicon	
10	Camera for gel electrophoresis with combs, power supply, transilluminator		

### **Exome enrichment of DNA libraries**

- We use SureSelect Baits v6 or v7 (Agilent) probes. They must be stored at -70  $^{\circ}$  C and carefully thawed on ice in the refrigerator beforehand!

- Hyb buffers #1,2,4 are stored on RT, cannot be cooled. Hyb 3 should be defrosted in advance.

- We use Blocking oligo: IGMblock11, IGMblock12. The stock concentration is 1000 µM.

#### **DNA library pooling**

Write down for each sample the concentration value (ng / mkl) measured by Qubit, select 10-13 libraries for 1 pool. In total, pool should weigh from 3 mkg to 5 mkg. The amount of each sample in the pool should be 350-500 ng (value may vary depending on the experimental goals).

Important: Make sure that the adapters in DNA libraries do not match.

Important: Libraries must be made with the same DNA insert size. You can pool libraries made with size - select x0.8 + x0.2, or libraries made with size - select x0.9 + x0.2.

It is necessary to completely dry the pool on the SpeedVac concentrator in 1.5 ml LoBind tubes at maximum vacuum and at temperature of 50°C.

#### Preparation of hybridization buffer

*Important: It* is necessary to observe the requirements for cleanliness of the workplace - before start you must wipe the work surface, pipettes and racks with 70% alcohol and leave all plastic (tips, tubes) in the box under the UV for 30 minutes.

1. Prepare Hybridization mix for 10 pools (the volume is such for uniform heating in the thermostat and get rid of the resulting precipitate) in 1.5 ml tube:

	x1	x10
- Hyb1 (20X SSPE)	9 µL	90 µL
- Hyb2(0.5M EDTA)	0.5 µL	5 µL
- Hyb3 (50X Denhardt's Solution)	3.5 µL	35 µL
- Hyb4 (10% SDS)	0.5 µL	5 µL

Vortex well and put in a thermostat at 65 ° C to get rid of the precipitate.

#### **Mixing Blocks and DNA library pools**

- 1. Add Blocking oligos to LoBind tube with dried pool of libraries:
  - Block 1 (cot DNA) =  $13 \,\mu L^*$

\*in case of reagent shortage you can reduce the volume to  $10 \ \mu L$ 

### Important! Pipet well and peel off the walls of the mix so that the dried DNA is FULLY DISSOLVED. Vortex the tubes so that the volume of the mixture is well mixed along the lower walls of the tube.

Transfer 13 µL pool and Block 1 mixtures to new 0.2 mL single PCR tubes. Sign LIBS.

Add 500 pmol to each pool of each Blocking oligo:

- IGMblock11 =	0.5 μL
- IGMblock12 =	0.5 μL
	=14 μL

2. Set the temperature program in the amplifier with a heated lid, the lid temperature should be 105 ° C. Put *Libs* in the thermocycler and start the program.

Temperature	Time	Actions
95°C	05:00	Put Libs
65°C	x	Put Hybs Put Probes

While the 95 ° C program is running, transfer 15  $\mu$ L of heated hybridization buffer into 0.2 ml tubes according to the number of pools, or it is allowed to transfer a sufficient volume for several reactions into one tube, sign *HYBS*. Put in the thermocycler for warming up as soon as the temperature reaches 65 ° C for at least 5 minutes.

#### **Probe preparation**

*Important:* Probes are of RNA nature, so the enrichment is performed with the most serious requirements to the cleanliness of the workplace, equipment and consumables.

- 1. Mix probes in single 0.2 mL tubes:
- Rnase Block =  $1 \mu L$
- SureSelect baits =  $4 \mu L$
- 2. Name the tubes **PROBES.** Place the tubes in a thermocycler at 65 ° C, allow to warm up for 4 minutes.

#### Hybridization of DNA probes and target areas

- 1. Wait until all components of the hybridization are heated up, the amplification program should switch to a constant mode of 65  $^{\circ}$  C.
- 2. Open tubes directly in the thermocycler and transfer 14 µ L Hybs to 14 µ L Libs , carefully mix by pipetting.
- 3. Then transfer 28 µ L of Hybs + Libs to 5 µ L Probes ( sum 33 µ L ), mix gently by pipetting and close the lids.
- 4. Important! Add mineral oil to prevent samples from evaporating; oil is added to the walls of the tubes so that it gently covers the reaction mixture. Close the lids tightly.
- 5. Leave hybridization for 18-24 hours;

Temperature	The time is .	Actions
		1. Transfer 14 μL Hybs to 14 μL Libs
65°C	œ	2. Transfer 28 μL Hybs+Libs to 5 μL Probes (sum = 33 μL).

#### Day 2. Washing the enrichment reaction products

This stage should be started in advance, approximately 1.5 hours before the end of hybridization.

#### **Prepare washing buffers**

A. Binding buffer

10 mL 5 M NaCl 500 uL 1 M Tris-HCl (pH 7.5) 100 uL 0.5 M EDTA 39.4 mL ddH2O

50 mL Total Volume

B. Wash Buffer #2

250 uL 20X SSC 500 uL 10% SDS 49.25 mL ddH2O

50 mL Total Volume

#### Washing the enrichment reaction products

Important! Place Dynabeads C1 on RT 30min before use.

Important! All stages of washing before elution and PCR should be carried out strictly at 65°C.

1. Prepare Wash Buffer 3 in 1.5 mL tube (per 1 pool) by mixing:

- Hyb4 =	12.1 μL
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- MQ Water =  $1200 \ \mu L$
- Wash Buffer  $2 = 300 \ \mu L$

Place the tube with wash buffer into the thermostat at 65°C for 45 minutes.

- 2. Take 30 μL Dynabeads C1 (per 1 pool) into 1.5 mL LoBind tube, place on a magnetic stand, wait until the solution becomes transparent, remove the supernatant;
- Remove the tube from the magnetic rack, add 200 µL Binding Buffer, Vortex+spin, place on the magnetic stand, wait 2 minutes until the solution becomes transparent, remove the supernatant;
- 4. Repeat step 3 twice to perform a total of 3 washes;

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5. Resuspend beads in 65  $\mu$ L Binding Buffer.

*Important:* You can wash the Dynabeads for several samples in one tube (up to 7 samples). To do this you need to increase the buffer volume in proportion to the number of samples. For example, for 4 reactions, take 120  $\mu$ L Dynabeads and wash them three times with 800  $\mu$ L Binding Buffer, then resuspend to 280  $\mu$ L Binding Buffer.

- 6. Add 5 µl/pool Block 2 (Salmon Sperm DNA 1 mg/µl) and place on an orbital shaker for 15 minutes at RT.
- 7. Transfer 70  $\mu$ L "locked" Dynabeads to new 1.5 mL LoBind tubes, sign according to the pools.
- 8. Put the tubes in the thermostat at 65 ° C, warm up for 5 minutes,
- 9. Open the thermocycler, transfer the entire volume of enriched pools to the Dynabeads in the thermostat, mix by pipetting;
- 10. Incubate the tubes at 65°C for 30 minutes in a thermoshaker;
- 11. Place the tubes on a magnetic stand, wait until the solution becomes transparent (~1 min), and the magnetic particles collect at the tube wall, remove the supernatant;
- 12. Add 500 µL of preheated Wash Buffer 3, Vortex+spin, put in a thermoshaker at 65°C for 10 minutes.
- 13. Carefully centrifuge the drops, transfer tubes to a magnetic stand, remove the supernatant;
- 14. Repeat steps 12-13 two more times (total 3 washes with WB3);
- 15. Dry beads on a magnetic stand for 3-4 minutes.
- 16. Resuspend in 31  $\mu L$  MQ.

#### **Post-capture PCR**

At this stage, enriched libraries are "detached" from probes during denaturation and then amplified.

**IMPORTANT: Before setting up the PCR, you must first "remove" = denaturate DNA from Dynabeads.** To do this, transfer the entire reaction volume (31  $\mu$ l) into the new 0.2  $\mu$ l PCR tubes, put in the thermocycler at 95 ° C for 5 minutes. Then quickly put on a magnetic stand and transfer the transparent supernatant into the new 1.5 mL LoBind tube.

In the PCR use half of the enriched sample (15-17  $\mu l),$  sign the 2nd half and store at -20  $^{\circ}$  C.

Use reagents from KAPA HiFi HotStart PCR Kit.

### Important: For PCR after enrichment the common reagent should NOT be used. Buffer, dNTPs and primer aliquots must be proprietary for WES only.

Primers used (at a concentration of  $30 \mu M$ ):

MGIAd_PCR_1	/5Phos/GAACGACATGGCTACGA
MGIAd_PCR_2	TGTGAGCCAAGGAGTTG

1. Mix the following components in a sterile PCR tube:

<ul> <li>MGI primer 2 =</li> <li>KAPA HiFi HotStart DNA Polymerase =</li> <li>DNA enriched fragments =</li> <li>mQ =</li> </ul>	10 μL 1.5 μL 8 μL 8 μL 1 μL 15-17 μL up to 50 μL
	Total: 50 $\mu$ L

1. Put into the thermocycler and start the following program:

Temperature	The time is .	Repeats
95°C	03:00	
98°C	00:20	
60°C	00:15	X10*
72°C	00:30	
72°C	10:00	
4°C	Storage	

\*NB! The number of PCR cycles may vary depending on the use of probes from different manufacturers. For 4µl Agilent, 10 cycles are optimal.

**2.** Select 2.5 μl of PCR product and immediately perform Gel-Electrophoresis (2% agarose gel, 30 min 130 V) for quality control. A good result is a length distribution with a maximum in the range of 300-500 bp.

Important! Step 3 is critical, if the gel is empty, it makes sense to reanimate the situation by adding a PCR mix and run few more PCR cycles.

3. If the gel is all OK, then proceed to further purification of the PCR product.

*Stopping point*: if processed the next day samples can be storeat +4°C overnight.

#### **Purification using Agencourt AMPure XP**)

- 1. Transfer the reaction to a new 1.5 ml LoBind tube
- 2. Add to 50  $\mu$ L reaction 50  $\mu$ L Ampure beads (1x of reaction volume), vortex+spin;
- 3. Incubate at room temperature for 7 minutes;
- 4. Transfe the tubes to a magnetic rack. After the mixture becomes transparent (it takes 7 minutes), you should carefully remove the supernatant without touching the beads that contain fragments of DNA;
- 5. Add 400 μL fresh (!) 80% EtOH, rotate the tube in a rack 180° so that the beads run through the EtOH solution several times, collect all the beads on one wall, gently remove EtOH without touching the bids;
- 6. Repeat step 5 one more time to get 2 washes;
- 7. Remove residues of 80% EtOH with a pipette with 2-20 tips. Leave the test tubes with an open lid to dry directly on the magnetic stand for 3-5 minutes, making sure that the beads do not dry out (when dried they lighten).
- 8. Add 38 μL mQ, resuspend beads well on a Vortex+Spin, incubate for 7 min, then place on a magnetic stand, incubate for 7 min. Transfer the supernatant to a new tube. The supernatant contains purified DNA.

#### **Enrichment quality control**

To carry out the quantitative control of the obtained enriched pools - to measure the concentration with the Qubit High Sense set according to the manufacturer's protocol.

The total amount of enriched pool should be at least 70 ng. If the amount is less, it is necessary to pool with other samples in the further reaction of circularization to obtain sufficient amount of DNA for circulation.

Optionally – perform quality control of the obtained enriched pools on the Agilent Bioanalyzer 2100 using the High Sensitivity Kit according to the manufacturer's protocol.

A good result is a length distribution with a peak of 250-400 bp. A good pool suitable for further preparation for sequencing is shown in *Figure 1*.

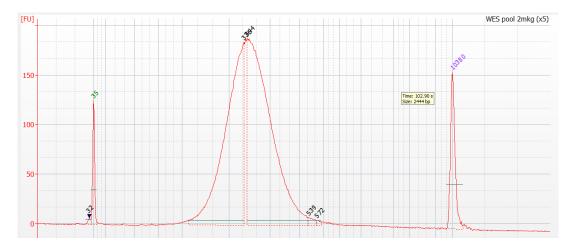


Fig.1. Chromatogram of enriched DNA library with 350 bp peak for MGI sequencing.