# Viability E12.5 Secondary Screen IMPC\_EVM \_001

#### **Purpose**

To assess the viability, sub-viability, and lethality of homozygous embryos at E12.5

## **Experimental Design**

- Set up timed matings with heterozygous mice
- Day 0 is defined as the midpoint of the prior dark cycle following the identification of a copulation plug.
- Collect embryos at E12.5
- Collect tissue and genotype embryos.

#### **Procedure**

- 1. Set up timed mating with heterozygous animals. Aim to dissect and collect >=28 alive embryos, otherwise lethal and subviable calls cannot be made. If more than three homozygous pups are produced before 28 pups are genotyped, a viable call can be made.
- 2. Collect tissue for genotyping and (OPTIONAL) score Gross Morphology and/or process for Histopathology and or Imaging.
- 3. Genotype all embryos and
  - a. Strains that produce NO existing homozygous embryos will be considered LETHAL (complete embryonic lethality [MP:TBC]).
  - b. Strains that produce NO live (absence of heartbeat) homozygous embryos will be considered LETHAL (complete embryonic lethality [MP:TBC]).
  - c. Strains that produce live homozygous embryos but with an obvious defect will be left to the discretion of the center with the decision and reason recorded in the parameters.
  - d. X-linked strains that produce NO live hemizygous male embryos from female carriers will be considered LETHAL (complete embryonic lethality [M P:TBC]).
- 4. Flag strains that produce less than normal numbers of homozygous/hemizygous male progeny
  - a. Strains that produce <50% expected homozygous progeny will be annotated as partial embryonic lethality [MP:TBC].
  - b. X-linked strains that produce <50% expected male hemizygous progeny from female carriers will be considered partial embryonic lethality [MP:TBC].

#### Notes

All genotypes should be collected using validated assays.

Y chromosome assay required for X-linked lethal strains.

#### Data Analysis, annotation and display (+statistics)

Preliminary: No analysis required as it is a line level procedure. This could change with additional data about the procedure

See E12.5 Gross Morphology protocol for MP calls of specific phenotypes at this time point.

Total Embryos: All, WT, Het, Hom

Alive, dead, and defect (all genotyped)

Total Dead: All, WT, Het, Hom

Total Defect (Alive or Dead): All, WT, Het, Hom

Abnormal and dead embryos
 Litter size: all genotyped embryos
 ignore partials and reabsorptions.

#### **Parameters and Metadata**

### Total live embryos IMPC\_EVM\_024\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

#### Total embryos IMPC\_EVM\_023\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: true Is Annotated: false

# Total live heterozygous IMPC\_EVM\_025\_001 | v1.0

simpleParameter

Req. Analysis: false	Req. Upload: false	Is Annotated: false		
Time of dark cycle start IMPC_EVM_020_001   v1.0 procedureMetadata				
Req. Analysis: false	Req. Upload: true	Is Annotated: false		
Outcome IMPC_EVM_	.001_001   v1.1			
simpleParameter				
Req. Analysis: false	Req. Upload: true	Is Annotated: true		
<b>Options:</b> Homozygous - Lethal, Hemizygous - Viable, Insufficient numbers to make a call, Hemizygous - Lethal, Homozygous - Subviable, Homozygous - Viable,				
Number of reabsorptions IMPC_EVM_015_001   v1.0 simpleParameter				
Req. Analysis: false	Req. Upload: false	Is Annotated: false		

## Total embryos WT IMPC\_EVM\_004\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: true Is Annotated: false

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## **% embryos WT** IMPC\_EVM\_016\_001 | v1.3

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

Unit Measured: %

**Derivation:** div('IMPC\_EVM\_004\_001', 'IMPC\_EVM\_023\_001')

#### Decision IMPC\_EVM\_002\_001 | v1.1

simpleParameter

Req. Analysis: false Req. Upload: true Is Annotated: false

**Options:** Go to E18.5, No further data available, Go to E14.5, Go to E14.5 and E18.5,

Go to E15.5, Go to E9.5,

#### Average Litter Size IMPC\_EVM\_019\_001 | v1.0

simpleParameter

Req. Analysis: false	Req. Upload: false	Is Annotated: false		
Time of dark cycle procedureMetadata	end IMPC_EVM_021_001	v1.0		
Req. Analysis: false	Req. Upload: true	Is Annotated: false		
Comment on Decision (in English) IMPC_EVM_003_001   v1.2 simpleParameter				
Req. Analysis: false	Req. Upload: false	Is Annotated: false		
Total gross defect PC_EVM_011_001   v1.2 simpleParameter	at dissection (alive	or dead) embryos IM		
Req. Analysis: false	Req. Upload: true	Is Annotated: false		
Total embryos heterozygous IMPC_EVM_005_001   v1.0 simpleParameter				
Req. Analysis: false	Req. Upload: true	Is Annotated: false		

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Embryo medium IMPC_EVM_022_001   v1.0				
procedureMetadata				
Req. Analysis: false	Req. Upload: true	Is Annotated: false		
Options: Warm PBS, Ice,				
Total embryos homozygous IMPC_EVM_006_001   v1.0 simpleParameter				
Req. Analysis: false	Req. Upload: true	Is Annotated: false		
Total live homozygous IMPC_EVM_027_001   v1.0 simpleParameter				
Req. Analysis: false	Req. Upload: false	Is Annotated: false		
Total live WT IMPC_EVM_026_001   v1.0 simpleParameter				
Req. Analysis: false	Req. Upload: false	Is Annotated: false		

% embryos homozygous IMPC\_EVM\_018\_001 | v1.3 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: false **Unit Measured:** % **Derivation:** div('IMPC EVM 006 001', 'IMPC EVM 023 001') Total gross defect at dissection (alive or dead) homozygous IMPC\_EVM\_014\_001 | v1.3 simpleParameter Req. Analysis: false Req. Upload: true Is Annotated: false Total dead embryos IMPC\_EVM\_007\_001 | v1.0 simpleParameter Req. Analysis: false Req. Upload: true Is Annotated: false

Total gross defect at dissection (alive or dead) heterozygous IMPC\_EVM\_013\_001 | v1.2

Req. Analysis: false	Req. Upload: true	Is Annotated: false		
Total dead homozy simpleParameter	<b>'gous</b> IMPC_EVM_010_00	01   v1.0		
Req. Analysis: false	Req. Upload: true	Is Annotated: false		
% embryos heterozygous IMPC_EVM_017_001   v1.3 simpleParameter				
Req. Analysis: false	Req. Upload: false	Is Annotated: false		
Unit Measured: %				
Derivation: div('IMPC_EVM_0	005_001', 'IMPC_EVM_023_00	01')		
Total dead WT IMPC_EVM_008_001   v1.0 simpleParameter				
Req. Analysis: false	Req. Upload: true	Is Annotated: false		

# Total gross defect at dissection (alive or dead) WT IMPC\_EV

M\_012\_001 | v1.2

simpleParameter

Req. Analysis: false Req. Upload: true Is Annotated: false

# Total dead heterozygous IMPC\_EVM\_009\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: true Is Annotated: false