Evaluation of the Invader® assay platform for molecular analysis of the Factor V (G1691A) and Factor II (G20210A) mutations.

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Aim:

- To evaluate the Third WaveTM Invader® DNA assay for the detection of the FVL (G1691A) and Factor II (G20210A) mutations.
- Test 100 samples where the genotype is known for the FV and FII mutations as determined by the RFLP method.
- Compare and contrast the two methods for use in a diagnostic setting.

Methodology:

The RFLP method (PCR based):

Factor V Leiden:

Restriction enzyme = Mnl I

(loss of restriction site)



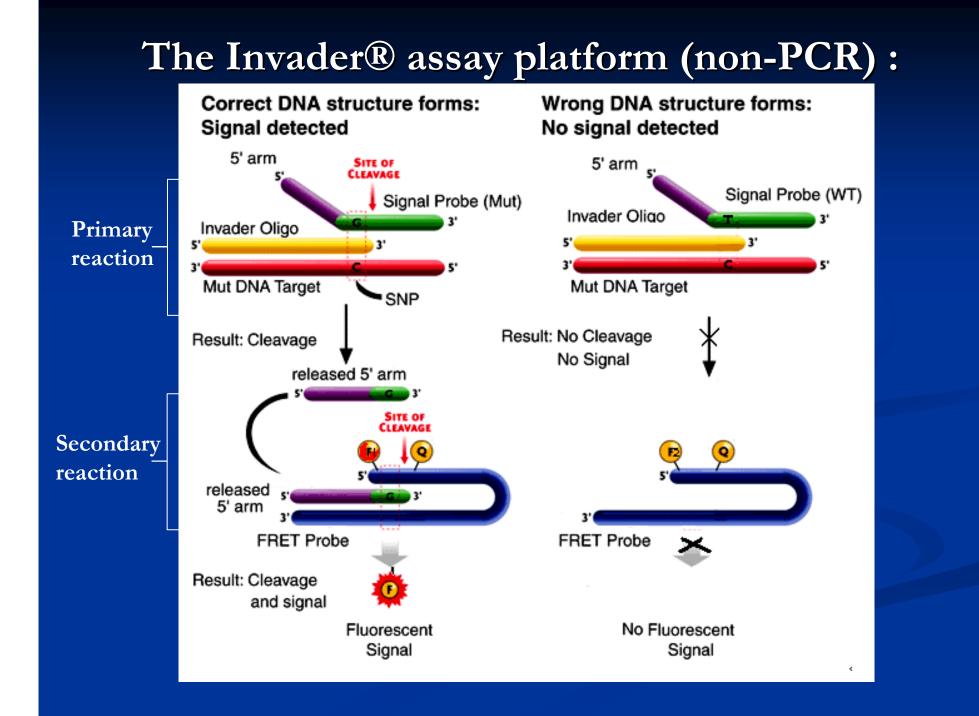
Undigested No DNA Normal Homozygous Heterozygous Factor II:

Restriction enzyme = Hind III

(Use of mutagenic primer and G > A creates restriction site)



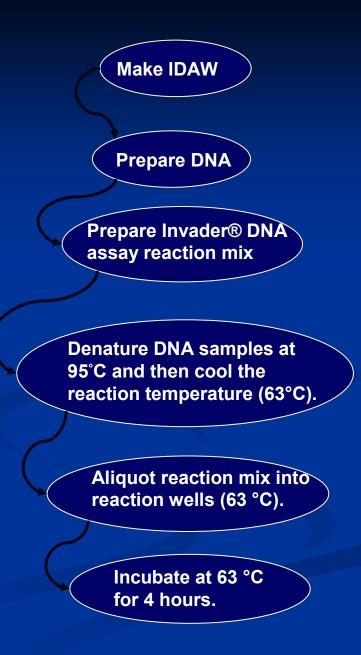
Undigested No DNA Heterozygous Normal



Invader® assay platform:

(Third Wave technologies)

- >100ng of DNA is needed
- 96 well plate format.
- biplex format F (WT) and R (Mut) signal
- Controls include a Normal, Heterozygote and Homozygote
- Fluorescence analysed using a 96 well plate reader (CytoFluor 96 well plate reader).
- An excel worksheet (IDAW) is used to calculate the net signal/ background or net fold over zero (FOZ).
- Ratio of the WT reaction to the mutant reaction.
 - Heterozygous = >0.3 to <3
 - Homozygous = <0.2
 - Normal = >5
- Total assay time = ~ 5 hours (hands on time = 30-45min)



Results:

IDAW:

| Data File Date Stamp: | | | | | | | | | | | |
|---|---|---|---|--|--|--|--|---|---|--|-----|
| F Signal | Date Stamp: | | | | | Raw Data | | | | | |
| (Mut) | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 |
| A | 259 684 | 211 | | | | | | | | | |
| B C | 909 | 285 232 | | | | | | | | | |
| Ď | 294 | 217 | | | | | | | | | |
| E | 349 318 | 199 212 | | | | | | | | | |
| F G | 1508 | 212 | | | | | | | | | |
| Ĥ | 352 | 417 | | | | | | | | | |
| | | | | | | | | | | | |
| R Signal (WT) | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 |
| A | 534 | 1678 | | | | | | | | | |
| B C | 540 245 | 227 2210 | | | | | | | | | |
| D | 253 | 911 | | | | | | | | | |
| E | 400 | 1026 | | | | | | | | | |
| F | 686 1374 | 1174 1495 | | | | | | | | | |
| G H | 1374 | 847 | | | | | | | | | |
| | | | | | | Lot Numbers | | | | FVL (1691) Control 1 (V | (T) |
| Operator: | Date: | DNA Reactio | n Buffer 1 (i | 30 | | | | ET R Cassette (R) | 0 | FVL (1691) Control 1 (M FVL (1691) Control 2 (H | |
| oly2 | | FVL (1691) I | | | | | | ET F Cassette (F) | | FVL (1691) Control 3 (M | |
| | | FVL (1691) I | | | | | | zyme 40 ng/µl (E) | | Control 4 (No Target Bla | |
| | | | Invader Data Analysis - FVL (G1691A) Biplex Assay | | | | | Version 040202 | | | |
| | Invader | | | | | Net F Signal | Net R Signal | | | | |
| Sample | Genotype | F Signal | R Signal | F Signal FOZ | R Signal FOZ | FOZ | FOZ | RATIO | Data | Action | |
| FVL (1691) | | | | | | | | | | | |
| Control 1 | WT | 259 | 534 | 0.88 | | | | | | | |
| | | 200 | | 0.00 | 2.11 | 0.04 | 1.11 | 27.767 | VALID | NONE | |
| EVI (1691) | | | | | | | | | | | |
| FVL (1691) Control 2 | HET | 684 | 540 | 2.33 | 2.11 | 1.33 | 1.11 | 0.855 | VALID | NONE | |
| Control 2 | | 684 | 540 | 2.33 | 2.13 | 1.33 | 1.13 | 0.855 | VALID | NONE | |
| Control 2 FVL (1691) | HET MUT | | | | | | | | | | |
| Control 2 FVL (1691) Control 3 | | 684 909 | 540 245 | 2.33 | 2.13 | 1.33 | 1.13 | 0.855 | VALID | NONE | |
| Control 2 FVL (1691) | | 684 | 540 | 2.33 | 2.13 | 1.33 | 1.13 | 0.855 | VALID | NONE | |
| Control 2 FVL (1691) Control 3 | MUT EQ1 | 684 909 | 540 245 | 2.33 | 2.13 0.97 1.58 | 1.33 | 0.04 | 0.855 | VALID VALID VALID INVALID | NONE NONE NONE REPEAT SAMPLE | |
| Control 2 FVL (1691) Control 3 Control 4 66(4ul) 6(4ul) | MUT EQ1 WT | 684 909 294 349 318 | 540 245 253 400 686 | 2.33 3.09 1.19 1.08 | 2.13 0.97 1.58 2.71 | 1.33 2.09 0.19 0.08 | 1.13 0.04 0.58 1.71 | 0.855 0.019 3.106 20.965 | VALID VALID VALID INVALID VALID | NONE NONE NONE REPEAT SAMPLE NONE | |
| Control 2 FVL (1691) Control 3 Control 4 66(4ul) | MUT EQ1 | 684 909 294 349 | 540 245 253 400 | 2.33 3.09 1.19 | 2.13 0.97 1.58 | 1.33 2.09 0.19 | 0.04 | 0.855 | VALID VALID VALID INVALID | NONE NONE NONE REPEAT SAMPLE | |
| Control 2 FVL (1691) Control 3 Control 4 66(4ul) 6(4ul) 9 10 | EQ1 WT HET WT | 684 909 294 349 318 | 540 245 253 400 686 1374 1302 | 2.33 3.09 1.19 1.08 5.13 1.20 | 2.13 0.97 1.58 2.71 5.43 5.15 | 1.33 2.09 0.19 0.08 4.13 0.20 | 1.13 0.04 0.58 1.71 4.43 4.15 | 0.855 0.019 3.106 20.965 1.073 21.017 | VALID VALID VALID INVALID VALID VALID VALID | NONE NONE NONE REPEAT SAMPLE NONE NONE NONE | |
| Control 2 FVL (1691) Control 3 Control 4 66(4ul) 6(4ul) 9 | MUT EQ1 WT HET | 684 909 294 349 318 1508 | 540 245 253 400 686 1374 | 2.33 3.09 1.19 1.08 5.13 | 2.13 0.97 1.58 2.71 5.43 | 1.33 2.09 0.19 0.08 4.13 | 1.13 0.04 0.58 1.71 4.43 | 0.855 0.019 3.106 20.965 1.073 | VALID VALID VALID INVALID VALID VALID | NONE NONE NONE REPEAT SAMPLE NONE NONE | |
| Control 2 FVL (1691) Control 3 Control 4 66(4ul) 6(4ul) 9 10 11 12 | EQ1 WT HET WT | 684 909 294 349 318 1508 352 | 540 245 253 400 686 1374 1302 | 2.33 3.09 1.19 1.08 5.13 1.20 | 2.13 0.97 1.58 2.71 5.43 5.15 | 1.33 2.09 0.19 0.08 4.13 0.20 | 1.13 0.04 0.58 1.71 4.43 4.15 5.63 0.04 | 0.855 0.019 3.106 20.965 1.073 21.017 | VALID VALID VALID INVALID VALID VALID VALID | NONE NONE NONE REPEAT SAMPLE NONE NONE NONE | |
| Control 2 FVL (1691) Control 3 Control 4 66(4ul) 6(4ul) 9 10 11 | EQ1 WT HET WT WT | 684 909 294 349 318 1508 352 211 | 540 245 253 400 686 1374 1302 1678 | 2.33 3.09 1.19 1.08 5.13 1.20 0.72 | 2.13 0.97 1.58 2.71 5.43 5.15 6.63 | 1.33 2.09 0.19 0.08 4.13 0.20 0.04 | 1.13 0.04 0.58 1.71 4.43 4.15 5.63 | 0.855 0.019 3.106 20.965 1.073 21.017 140.810 | VALID VALID VALID INVALID VALID VALID VALID VALID | NONE NONE NONE REPEAT SAMPLE NONE NONE NONE | |
| Control 2 FVL (1691) Control 3 Control 4 66(4ul) 6(4ul) 9 10 11 12 | MUT EQ1 WT HET WT WT | 684 909 294 349 318 1508 352 211 285 | 540 245 253 400 686 1374 1302 1678 227 | 2.33 3.09 1.19 1.08 5.13 1.20 0.72 0.97 | 2.13 0.97 1.58 2.71 5.43 5.15 6.63 0.90 | 1.33 2.09 0.19 0.08 4.13 0.20 0.04 0.04 | 1.13 0.04 0.58 1.71 4.43 4.15 5.63 0.04 | 0.855 0.019 3.106 20.965 1.073 21.017 140.810 1.000 | VALID VALID VALID INVALID VALID VALID VALID VALID INVALID | NONE NONE REPEAT SAMPLE NONE NONE NONE REPEAT SAMPLE | |
| Control 2 FVL (1691) Control 3 Control 4 66(4ul) 6(4ul) 9 10 11 12 13 | MUT EQ1 WT HET WT WT WT | 684 909 294 349 318 1508 352 211 285 232 | 540 245 253 400 686 1374 1302 1678 227 2210 | 2.33 3.09 1.19 1.08 5.13 1.20 0.72 0.97 0.79 | 2.13 0.97 1.58 2.71 5.43 5.15 6.63 0.90 8.74 | 1.33 2.09 0.19 0.08 4.13 0.20 0.04 0.04 0.04 | 1.13 0.04 0.58 1.71 4.43 4.15 5.63 0.04 7.74 | 0.855 0.019 3.106 20.965 1.073 21.017 140.810 1.000 193.379 | VALID VALID VALID INVALID VALID VALID VALID VALID VALID VALID | NONE NONE NONE REPEAT SAMPLE NONE NONE NONE REPEAT SAMPLE NONE | |
| Control 2 FVL (1691) Control 3 Control 4 66(4ul) 6(4ul) 9 10 11 12 13 19 | MUT EQ1 WT HET WT WT WT WT | 684 909 294 349 318 1508 352 211 285 232 217 | 540 245 253 400 686 1374 1302 1678 227 2210 911 | 2.33 3.09 1.19 1.08 5.13 1.20 0.72 0.97 0.79 0.74 | 2.13 0.97 1.58 2.71 5.43 5.15 6.63 0.90 8.74 3.60 | 1.33 2.09 0.19 0.08 4.13 0.20 0.04 0.04 0.04 0.04 | 1.13 0.04 0.58 1.71 4.43 4.15 5.63 0.04 7.74 2.60 | 0.855 0.019 3.106 20.965 1.073 21.017 140.810 1.000 193.379 65.020 | VALID VALID VALID INVALID VALID VALID VALID VALID VALID VALID VALID | NONE NONE NONE REPEAT SAMPLE NONE NONE NONE REPEAT SAMPLE NONE NONE | |
| Control 2 FVL (1691) Control 3 Control 4 66(4ul) 6(4ul) 9 10 11 12 13 19 15 | EQ1 WT HET WT WT WT WT WT | 684 909 294 349 318 1508 352 211 285 232 217 199 | 540 245 253 400 686 1374 1302 1678 227 2210 911 1026 | 2.33 3.09 1.19 1.08 5.13 1.20 0.72 0.97 0.79 0.74 0.68 | 2.13 0.97 1.58 2.71 5.43 5.15 6.63 0.90 8.74 3.60 4.06 | 1.33 2.09 0.19 0.08 4.13 0.20 0.04 0.04 0.04 0.04 0.04 0.04 | 1.13 0.04 0.58 1.71 4.43 4.15 5.63 0.04 7.74 2.60 3.06 | 0.855 0.019 3.106 20.965 1.073 21.017 140.810 1.000 193.379 65.020 76.383 | VALID VALID VALID INVALID VALID VALID VALID VALID VALID VALID VALID VALID VALID | NONE NONE NONE REPEAT SAMPLE NONE NONE NONE REPEAT SAMPLE NONE NONE NONE | |

Factor V and Factor II Invader results:

| | Factor V | Factor II |
|-------------------|----------|-----------|
| Number of samples | 110 | 110 |
| Normal | 71 | 98 |
| Heterozygous | 33 | 9 |
| Homozygous | 2 | 0 |
| Fails | 4 (3.6%) | 3 (2.7%) |
| % Repeated | 5.5% | 3.6% |
| % Concordance | 100% | 100% |

Discussion:

The genotypes obtained from the Invader® assay showed 100% concordance to the RFLP method showing that it is suitable for use in diagnostic molecular genetics.

<u>Repeated samples:</u> Factor V 5.5% and Factor II 3.6%.

Possible reasons:

- Low signal as a result of low DNA concentration meaning the patient sample doesn't exceed the background fluorescence seen in the no target blank control.
 - when repeated with more DNA the correct genotype was obtained.
- Failed samples: Factor V 3.6% and Factor II 2.7%.
- The samples that could not be genotyped were due to a low DNA concentration.

- 50 /110 samples tested were from Southampton Human genetics unit (total volume 10-15µl).
- All failed samples were from this source and could not be quantified or genotyped due to sample being depleted.

Comparison of Invader and RFLP:

| | RFLP | Invader platform |
|-----------------|------------------------------|----------------------------|
| Total time | 7-25 hours | 4-5 hours |
| Hands on time | 2-3 hours | <45 mins |
| Number of steps | 8 | 4 |
| | 1: Prepare sample | 1: Prepare sample |
| | 2: Make master mix | 2: Make master mix |
| | 3: Make digest mix | 3: Add master mix to plate |
| | 4: Add digest mix to samples | 4: Read plate |
| | 5: Restriction digest | |
| | 6: Pour gel | |
| | 7: Load gel | |
| | 8: Image gel | |
| Analysis | Gel (subjectivity) | Excel spreadsheet (IDAW) |
| Visualisation | EtBr (mutagenic) | FRET (fluorescence) |

Cost per test:

Invader assay platform = $\pounds 8$ per result

 $RFLP = Approx. \pounds 2-3$

However, the Invader assay requires the use of a flourescence plate reader which costs around $\pounds 9000$

Other applications of the Invader assay platform:

- Cystic fibrosis testing
- Connexin 26
- MTHFR (methylenetetrahydrofolate reductase)
- ApoE
- Hexosaminidase A (Tay Sachs)

Conclusion:

- The invader offers cheap, rapid detection of SNP's
- Highly reproducible results with 100% concordance to existing methods.
- It is non-PCR based.
- Can test up to 92 samples at once
- Wide applications to other areas of molecular diagnostics.
- Highly dependant on template concentration as it affects the reaction dynamics and overall signal strength. However, 120/120 tests on DNA from our lab worked first time.
- More expensive than the existing RFLP method but is less labour intensive and is more rapid with results in ~ 5 hours.