ConfirmPCR™ assays

Our best-in-class ConfirmPCR™ assays covering all human canonical exons of protein coding genes are highly versatile and can be used in any workflow.

WGS/WES variant confirmation

1 million ConfirmPCR™ assays, of which over 10,000 have been validated in variant confirmation studies

NGS gap filling

Extend your preferred enrichment method with our ConfirmPCR™ assays to enrich low coverage regions

Targeted resequencing

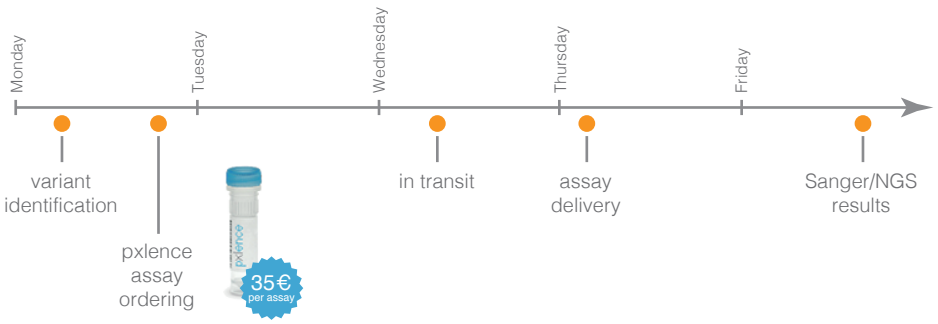
54 validated gene panels totalling up to 223 genes and 4008 ConfirmPCR™ assays

All our ConfirmPCR™ assays have been thoroughly validated *in silico*, resulting in an off-the-shelf PCR success rate of >97%. The optimized design parameters allow universal PCR conditions while ensuring uniform sequencing coverage.

- ✓ Avoid SNPs in primer annealing sites
- ✓ Minimize secondary structures
- ✓ Maximize primer specificity
- ✓ Optimize GC content, annealing temperature and other parameters

From variant identification to confirmation in one week

Save time and money on logistics, administration, assay design and variant validation by using pxlence's ConfirmPCR™ assays.



Contact

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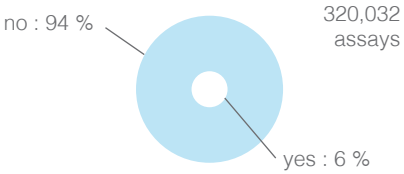
Best-in-class PCR assay quality

To showcase the quality of our ConfirmPCR™ assays, we sequenced a random set of 2300 different amplcons. The results show that 95% of the assays generate more than 98% on-target specificity. This correlates very well with the *in silico* score depicted on our website, allowing you to critically assess the quality of each of our assays. More details about the assay quality characteristics can be found in our Biomolecular Detection and Quantification paper (*PubMed ID 27077044*).



SNP present in primer annealing sites

ConfirmPCR™ assay coverage specificity

Some of our customers

Dr. Hans Atli Dahl
CEO of Amplexa Genetics, Odense, Denmark

> *At Amplexa Genetics we have been using pxlence assays for some time now and we are very satisfied with the solution. It is saving us valuable time at no extra cost. Prior to pxlence we spent a lot of time on designing primers and subsequent testing and optimizing the amplicons. Now we just order the pre-designed and ready-to-use amplicons from pxlence and run them at the defined standard conditions – works every time. Adds no extra cost but saves us a lot of time!*

References

Our ConfirmPCR™ assays have been used in the development of a wide range of NGS gene panels, both for research and diagnostic purposes. In addition, our customers use the assays for Sanger-based validation of variants.

Targeted resequencing and variant validation using pxlence PCR assays
2294 assays - PubMed ID 27077044

Massively parallel sequencing for early molecular diagnosis in Leber congenital amaurosis
16 genes - 375 assays - PubMed ID 22261762

Molecular diagnostics for congenital hearing loss including 15 deafness genes using a next generation sequencing plaform
16 genes - 376 assays - PubMed ID 22607986

Flexible, scalable, and efficient targeted resequencing on a benchtop sequencer for variant detection in clinical practice
265 genes - over 4000 assays - PubMed ID 25504618

More papers using our assays are available on our website.