# Development of a new LAMP assay for diagnosing the main meningitis pathogens

## Didia AMF<sup>1</sup>, Diallo K<sup>1,2,3\*</sup>, Amoikon TLS<sup>1</sup>, Tuo KJ<sup>1,4</sup>, Missa K.F<sup>1,5</sup>, Harrison OB<sup>3</sup>, Maiden MCJ<sup>3</sup>

1. Centre Suisse de Recherches Scientifiques en Côte d'Ivoire (CSRS), Abidjan, Côte d'Ivoire; 2. West African Centre for Cell Biology of Infectious Pathogens (WACCBIP), Accra, Ghana; 3. Department of Biology, University of Oxford, UK, 4. Institut National Polytechnique Felix Houphouët-Boigny, Yamoussoukro, Côte d'Ivoire; 5. Université Felix Houphouët-Boigny, Abidjan, Côte d'Ivoire



## Introduction

Meningitis is a serious disease with significant burden in low- and middle-income countries. Current diagnostic tests are expensive and difficult to use in secondary health centres where resources are limited. LAMP assays could be an alternative method to bring molecular detection closer to patients. This study therefore aimed to develop standardized LAMP assays that could detect in parallel reactions the four main pathogens causing meningitis using improved gene targets

## Methods

- Three LAMP reaction conditions were tested to determine the best method : the method of McKenna et al., 2011, the method of Tanner et Evans, 2014 and the method of Kim et al., 2012;
- ✓ Optimisation tests were carried out for the LAMP condition, which did not work in the standard condition, by varying the temperature and by varying the concentration of DNA polymerase.
- ✓ Following the tests, the LAMP condition described by Kim et al., 2012 was chosen as the best performing.
- ✓ The samples were subjected to the LAMP test and the results were analysed. The performance of the test was evaluated using various formulae, including sensitivity, specificity, negative predictive value and positive predictive value.



**Figure 1:** Flow diagram of LAMP test for detection of *Haemophilus influenzae*, Group b *Streptococcus*, *Neisseria meningitidis* and *Streptococcus pneumoniae* 

Target	FP	FN	ТР	TN	Sensitivity (%)	Specificity (%)	PPV (%)	VPN (%)
fucK	0	0	7	42	100	100	100	100
sodC	0	0	13	36	100	100	100	100
cfb	0	1	6	42	86	100	100	97
psaA	0	0	8	41	100	100	100	100

FP : False Positive, FN : False Negative, TP : True Positive, TN : True Negative, PPV : Positive predictive value, NPV : negative predictive value

Table 2: LAMP assay statistics

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Target	Name	Séquence
Fuck	F3-fucK	GATGTTTTCCAAAAATGGCTAA
	B3-fucK	TCCAACTTTTTCACCTGCA
	FIP-fuck	CCATTGTGTGATCTGTAGTGAATTGCGTTTTTATTTCGTCA
		ATGCT
	BIP-fuck	CGGGAACATCAATGATGACAAACATGGTTATTACTTAAAC
		CCAGC
	LB-Fuck	TTGGGATCCATCGATTTTAGCATC
Cfb	F3-Cfb	GGTGCATTGTTATTTTCACCA
	B3-Cfb	TCAACACTAGTAATAGCCTCA
	FIP-Cfb	GCCATTTGCTGGGCTTGATTATTACTTTTAGTACATGCTGA
		TCAAGTGAC
	BIP-Cfb	AGCTTGATCAAGATAGCATTCAGTTTTTTTAACCGGTTTT
		TCATAATCTGTTC
	LF-Cfb	ACATGATTTACCACTTGTGGAG
Sod C	F3-Sod C	GTAACAAAGATGTGGGTACAG
	B3-Sod C	CCATGGGTAACCATGTTGT
	FIP-Sod C	GCCTTCGCTTAATCCTTGTAAATCATTTTTGACTATTACTC
		AATCTAACTATGG
	BIP-Sod C	CCAAGCTGTGAGCCAAAAGATTTTCACCTTTAGGATCCC
		AGTG
	LB-Sod C	GACAGCTGGTTTAGGCGC
PsaA	F3- PsaA	CCAAGTGCCTACATCTGG
	B3-PsaA	AATCATAGCGATGACGACTA
	FIP-PsaA	GCAAGTAATGTTGCTCATTTTCTGATTTTATTGCATATTGC
		CAGTGC
	BIP-PsaA	CCTTCCAACGGAACCGGTATTTTTCGAGTACTCAAACGG
		AGTTC
	LB-PsaA	GAACAAGTGGCTTATTTGGATAACG

**Table 1:** Target primers designed for the detection of Haemophilus influenzae,Group b Streptococcus, Neisseria meningitidis and Streptococcus pneumoniae bylamp assay

## Results

✓ Of all the targets tested, only SodC, psaA, cfb and fucK showed

#### A B C D E F G H I J K L M N O A B C D E F G H I J K L M N O \* A B C D E F G H I J K L M N O \* A B C D E F G H I J K L M N O

signs of amplification using all three reaction conditions. However, the LAMP reactions using Thermopol buffer and the Isotherm were not reproducible. These reaction conditions were therefore abandoned. The final reaction using 1X housemade buffer worked well, was reproducible and therefore was selected to perform the validation assays.

- ✓ The LAMP test targeting the *fucK* gene showed a sensitivity, specificity, positive predictive value and negative predictive value of 100%. All *Haemophilus influenzae* samples tested positive, while the other bacteria in the panel tested negative (Table 2; figure 2).
- ✓ The LAMP test targeting the cfb gene revealed that 6 out of 7 Streptococcus agalactiae were detected by the test, while the other bacteria in the panel were negative. Sensitivity was therefore 86%, specificity and positive predictive value 100% and negative predictive value 97% (Table2; figure 3).
- ✓ The LAMP test targeting the sodC gene showed a sensitivity, specificity, positive predictive value and negative predictive value of 100%. All Neisseria meningitis samples tested positive, while the other bacteria in the panel tested negative (Table 2;figure 4).
- ✓ The LAMP test targeting the *psaA* gene showed a sensitivity, specificity, positive predictive value and negative predictive value of 100%. All *Streptococcus pneumoniae* samples tested positive, while the other bacteria in the panel tested negative (Table 2;figure 5).



Figure 2: Visualisation of products resulting from the LAMP assay targeting the *H. influenzae fucK* gene

#### A B C D E F GHI J K L M N O A B C D E F G H I J K L M N O \* A B C D E F G H I J K L M N O



Figure 3: Visualisation of products resulting from the LAMP assay targeting the S. agalactiae cfb gene



Figure 4: Visualisation of products resulting from the LAMP assay targeting the *N. meningitidis sodC* gene

#### A B C D E F G H I J K L M N O A B C D E F G H I J K L M N O \*ABCDEFG H I J K L M N O



Figure 5: Visualisation of products resulting from the LAMP assay targeting the S. pneumoniae psaA gene

# Conclusion

we have successfully developed rapid and sensitive LAMP assays for the detection of *H. influenzae, S. agalactiae, N. meningitidis*, and *S. pneumoniae.* They have been validated on DNA extracted from bacterial strains. Further validation with clinical samples is needed alongside further development to decrease the amount of sample handling required, making it more user friendly, and ensuring and easier uptake in peripheral settings. Assays using methodologies like LAMP holds great promise for improving meningitis diagnosis and could contribute to more effective management and treatment of the disease in resource limited settings.

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