



Development of high yielding, early maturing *Grain Amaranth*

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Introduction

Leafy vegetables of *Amaranthus* are a significant dietary component in Sub-Saharan Africa, with black and pale-seeded varieties being the most popular. Globally, pale-seeded (grain) *Amaranthus* holds great potential due to its high protein content, making it suitable for confectioneries, biscuits and pastries. However, its production and utilization are limited, leaving it underutilized and lacking commercial recognition despite its immense nutritional and economic potential. This study seeks to enhance the production levels of *Amaranthus* for both leaf and seed to address these challenges.

Materials & Methods

Grain amaranth seeds were selected from advanced populations derived from breeding NHAMARI. The following procedures were employed:

1. DNA Extraction: SDS extraction protocol with modifications to improve DNA quality using the dried bead method.
2. Quality Analysis: DNA quality was assessed through agarose gel electrophoresis.
3. Sequencing: Amplified fragments were sequenced using an Applied Biosystems 3130xl Genetic Analyzer and the BigDye terminator v3.1 cycle sequencing kit.
4. Genetic Analysis: BioEdit software and MEGA X were used for nucleotide analysis and phylogenetic assessments.



Fig. 1: Cream Seed Amaranth



Fig. 2: Seed Amaranth

Conclusion

The genetic improvement and field trials of cream-seeded *Amaranthus* have successfully produced a new high-yielding, early-maturing variety. Its enhanced yield, nutritional potential, and excellent acceptability make it a promising candidate for addressing the underutilization of grain *Amaranthus* and fostering its commercial recognition.

Results

Field Performance: The cream-seeded *Amaranthus* demonstrated excellent performance in farmers' fields, achieving an overall acceptability score of 88.5 out of 100.

Growth Characteristics: The first leaf harvest was achieved 10 days after transplanting.

Genetic Differentiation: Sequencing identified and separated NHAMARI from the cream-seeded *Amaranthus*, grouping them into two distinct phylogenetic clusters.

Yield: The new variety outperformed the previously released NHAMOLA5 (NHAMARI), yielding 2.8 t/ha compared to 2.5 t/ha for NHAMARI.

>NHAMARI

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CTTAATSTAAGTGCCAAGATTAACAATAAGACTTATTACTC
CTGATTATGAAACCCTARATACTGATATCTTGGCAGCATTCC
GAGTAAGTCCTCAACCTGGAGTCCACCTGAAGAAGCGGG
GGCTGCAGTAGCTGCCGAATCTTCTACTGGTACATGGACAA
GTGTATGGACCGACGGACTTACCAATCTTGATCGTTACAAA
GGACGATGCTACAACATCGAGCCCGTTGCTGGAGAAGAAA
ATCAATATATTTGTTATGTAGCGTATCCTTTAGACCTTTTGA
AGAAGGTTCTGTTACTAACATGTTTACTTCCATTGTGGGKAA
CGYATTTGGGTTCAAARCTTTGCGTGCTCTACMTTGGGAAG
ATTTGCGAATCCCTGTGCTTATGTCAAACTTTCCAAGGSC
CKCCTCWCGGWATCCAGGGTGAAGARATAAATTGAACA
AGYACGGSCGTCCCCTATTGGGATGCACTATTAACCASAAT
TGGGGGGATCCGCTAAAAACTAT
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>AMARANTHUS CREDM SEED

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AKYYYSSGTTKCWYMAGAYATRMSRMTAWGRCTKMKKAK
ACAYYYKRWKTWWSAAAACCTAGATACTGATATCTTGGC
AGCATTCCGAGTAAGTCTCAACCTGGAGTCCACCTGAAG
AAGCGGGGGCTGCAGTAGCTGCCGAATCTTCTACTGGTACA
TGGACAAGTGTATGGACCGACGGACTTACCAATCTTGATCG
TTACAAAGGACGATGCTACAACATCGAGCCCGTTGCTGGA
GAAGAAAATCAATATATTTGTTATGTAGCGTATCCTTTAGACC
TTTTTGAAGAAGGTTCTGTTACTAACATGTTTACTTCCATTG
TGGGTAACGTATTTGGGTTCAAAGCTTTGCGTGCTCTACGT
TTGGAAGATTTGCGAATCCCTGTTGCTTATGTCAAACTTTC
CAAGGCCCGCTCACGGTATCCAGGTTGAAAGAGATAAATT
GAACAAGTACGGTCTGCCCTATTGGGATGCACTATTAAC
CAAAATTGGGGKTATCCGCTAAAAACTATGGTCGAGCATGT
TATGAATGTCTTCGCGGTGGACTTGATTTTACAAM
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>NHOKRA1

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ATTMCGTCCCAAACAGAGACTAAAGYGC GGWGGACTTGA
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Sample Codes	Identity
NHAMARI	<i>Amaranthus hypochondriacus</i>
AMARANTHUS CREDM SEED	<i>Amaranthus retroflexus</i>