

*Article*

# **Phytochemical Analysis and Antioxidant, Antimicrobial, and Antiaging Activities of Ethanolic Seed Extracts of Four *Mucuna* Species**

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**Supplementary material**

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                  10                  20                  30                  40                  50  
**MGG** ATGCGATACT TGGTGTGAAT TGCAGAATCC CGTGAACCAT CGAGTCTTTG  
**MMM** ATGCGATACT TGGTGTGAAT TGCAGAATCC CGTGAACCAT CGAGTCTTTG  
**MIT** ATGCGATACT TGGTGTGAAT TGCAGAATCC CGTGAACCAT CGAGTCTTTG  
**MPR** TTGCGATACT TGGTGTGAAT TGCAGAATCC CGTGAACCAT CGAGTCTTTG

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                  60                  70                  80                  90                  100  
**MGG** AACGCAAGTT GCGCCCCAAG CCATTAGGTT GAGGGCACGC CTGCCTGGGT  
**MMM** AACGCAAGTT GCGCCCCAAG CCATTAGGTT GAGGGCACGC CTGCCTGGGT  
**MIT** AACGCAAGTT GCGCCCCAAG CCATTAGGTT GAGGGCACGC CTGCCTGGGT  
**MPR** AACGCAAGTT GCGCCCCAAG CCATTAGGCC GAGGGCACGC CTGCCTGGGT

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                  110                  120                  130                  140                  150  
**MGG** GTCACACATC GTTACCCTAA AGY-AAACGT CTCATGTGCG TTTGCAGGGT  
**MMM** GTCACACATC GTTACCCTAA AGCAAAACGT CTCATGTGCG TGTGCAGGGT  
**MIT** GTCACACATC GTTACCCTAA AGC-AAACGC CTCATGCGCG TGTGCAGGGT  
**MPR** GTCACACATC GTTACCCTAA ATGCAAACGC CTCACGTGCG TGCAGAGGGT

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                  160                  170                  180                  190                  200  
**MGG** GGAAGTTGAC CTCCCGTGGG CCACAA--CT CGCGGCTGGT TAAAAATGCA  
**MMM** GGAAGCTGAC CTCCCGTGGG GCACGACTCT CGCGGCTGGT TGAAAAATGGA  
**MIT** GGAAGCTGAC CTCCCGTGGG CCACGA--CT CGCGGCTGGT TGAAAAATGGA  
**MPR** GGATGCTGAC CTCCCGCGAG CATCGT--CT CGTGGCTGGT TGAAAAATCGA

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                  210                  220                  230                  240                  250  
**MGG** GTTCATGGTT GAGAATGCCG TGATAAAATG GTGGATGAGC ATTGCTCGAG  
**MMM** GTTCACGGTT GAGAATGCCG TGATAAAATG GTGGATGAGC GTTGCTCGAG  
**MIT** GTTCATGGTT GAGAATGCCG TGATAAAATG GTGGATGAGC ATTGCTCGAG  
**MPR** GTCCGCGGCC GAGCTCGTCG CGACAAAATG GTGGATGAGC GATGCTCGAG

.....|.....|.....|.....|.....|.....|  
                  260                  270                  280                  290                  300  
**MGG** ACCAATCGCG TGCTACTCAG TTAATTTTGG ACTCCTTGAC CCAKAW-GCA  
**MMM** ACCAATCGCG TGCTACTCAG TTAATTTTGG ACTCTTTGAC CCAGAT-GCG  
**MIT** ACCAATCGCG TGCGACTCGG TCAATTTTGG ACTCTTCGAC CCAAATCGCG  
**MPR** ACCAGTCCGG CCGGACCCGG CCAAGGTCGG ACTCCCCGAC CCTACACGGC

.....|.....|.....|.....|.....|.....|  
                  310                  320                  330                  340                  350  
**MGG** TCCTCGGATG CTCCCAACGA GACCTCAGGT CAGGCGGGGC CACCCGCTGA  
**MMM** TCCTCGGACG CTCCCAACGA GACCTCAGGT CAGGCGGGGC TACCCGCTGA  
**MIT** TCCACGGACG CTCCCAACGA GACCTCAGGT CAGGCGGGGC CACCCGCTGA  
**MPR** TCCACGGACG CTCCCAACGA GACCTCAGGT CAGGCGGGGC TACCCGCTGA

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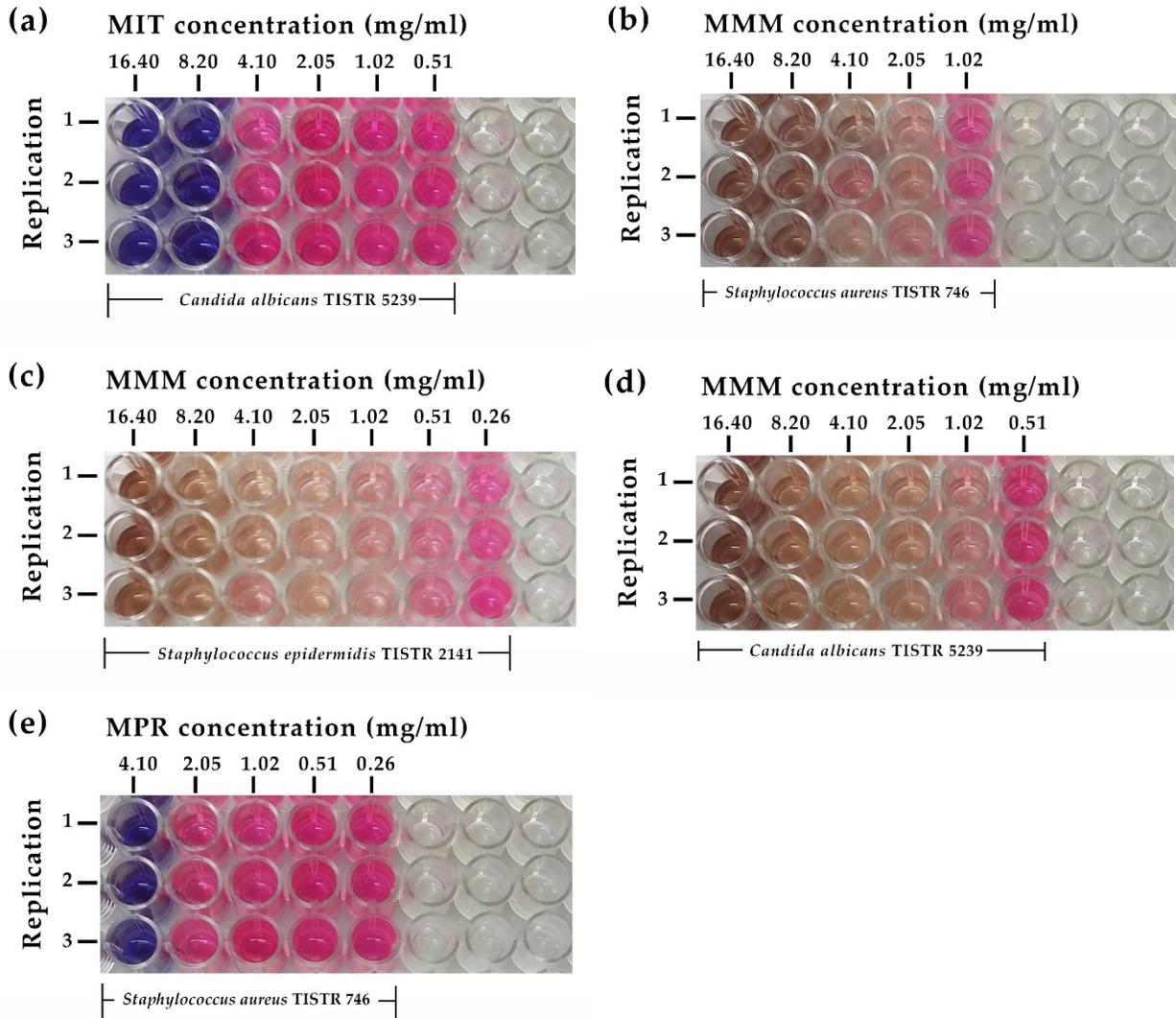
      .....|.....| .....|.....| .....|.....| .....|.....| .....|.....|
            360          370          380          390          400
MGG      GTTTAAGCAT ATCAATAAGC GGAGGAAAAG AACTAACAA GGATTCCCTT
MMM      GTTTAAGCAT ATCAATAAGC GGAGGAAAAG AACTAACAA GGATTCCCTT
MIT      GTTTAAGCAT ATCAATAAGC GGAGGAAAAG AACTAACAA GGATTCCCTT
MPR      GTTTAAGCAT ATCAATAAGC GGAGGAAAAG AACTAACAA GGATTCCCTT

      .....|.....| .....|.....| .....|.....| .....|.....| .....|.....|
            410          420          430          440          450
MGG      AGTAACGGCG AGCGAACCGG GAAGAGCCCA CCATGAAAAT CGGTCGTCAT
MMM      AGTAACGGCG AGCGAACCGG GAAGAGCCCA CCATGAAAAT CGGTCGTCAT
MIT      AGTAACGGCG AGCGAACCGG GAAGAGCCCA CCATGAAAAT CGGTCGTCAT
MPR      AGTAACGGCG AGCGAACCGG GAAGAGCCCA CCATGAAAAT CGGTCGCCCT

      .....|.....| .....|.....| .
            460          470
MGG      CGGCGTCCGA ATTGTAGTCT G
MMM      CGGCGTCCGA ATTGTAGTCT G
MIT      CGGCGTCCGA ATAGTAGTCT G
MPR      CGGCGTCCGA ATTGTAGTCT G

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**Figure S1:** Alignment of nucleotide sequences of the ribosomal DNA ITS2 regions of four *Mucuna* species: *M. gigantea* (MGG), *M. interrupta* (MIT), *M. monosperma* (MMM), and *M. pruriens* (MPR) (continued)



**Figure S2:** Minimal inhibitory concentrations (MICs) of *Mucuna* seed extracts on skin-related pathogen evaluated by broth micro-dilution and resazurin reduction methods. (a) Resazurin reduction of *Candida albicans* TISTR 5239 incubated with *M. interrupta* (MIT), (b) Resazurin reduction of *Staphylococcus aureus* TISTR 746 incubated with *M. monosperma* (MMM), (c) Resazurin reduction of *Staphylococcus epidermidis* TISTR 2141 incubated with *M. monosperma* (MMM), (d) Resazurin reduction of *Candida albicans* TISTR 5239 incubated with *M. monosperma* (MMM), and (e) Resazurin reduction of *Staphylococcus aureus* TISTR 746 incubated with *M. pruriens* (MPR). MICs were measured at the lowest concentrations of tests that did not change to pink color.

**Table S1:** Phytochemical constituents of *Mucuna* seed extracts evaluated by GC-MS analysis.

No.	MGG			MIT			MMM			MPR		
	RT	Compounds	%Area	RT	Compounds	%Area	RT	Compounds	%Area	RT	Compounds	%Area
1	6.79	Azetidine	8.3	6.01	Undecane	18.5	6.01	Undecane	3.2	6.07	1,2,3-Propanetriol, monoacetate	13.6
2	12.36	2,4-Di-tert-butylphenol, acetate	66.7	19.78	2-amino-6-cholo-3-cyano-5-(3-methyl-1-butenyl) pyrazine	21.5	6.29	Maltol	9.7	8.07	Butyl 2-methylbutanoate	11.6
3	16.86	2-Propenenitrile, 2-chloro-	8.3	21.65	Bis(2-ethylhexyl) phthalate	43.1	11.07	Cytidine	2.5	16.49	Palmitic acid	27.9
4	18.77	Urea, methyl-	16.7	23.39	Hexamethylcyclotrisiloxane	16.9	14.47	Resorcinol	9.0	18.15	Linoleic acid	29.3
5	-	-	-	-	-	-	16.48	Palmitic acid	13.2	18.38	Stearic acid	8.8
6	-	-	-	-	-	-	18.19	Dodecanamide, N-(2-hydroxyethyl)	29.8	19.74	4-Hydroxy-2,2,6-trimethylcyclohex-2-enone	4.8
7	-	-	-	-	-	-	19.78	5-(N-piperidyl)-4-oxopentanoic acid piperidide	9.1	21.14	2-Ethylacridine	2.0
8	-	-	-	-	-	-	21.66	Bis(2-ethylhexyl) phthalate	23.5	21.66	Bis(2-ethylhexyl) phthalate	2.0