

- pyaudi: A Taylor polynomial algebra toolbox for
- differentiable intelligence, automatic differentiation,
- and verified integration applications.
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Software

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Summary

pyaudi is a Python toolbox developed at the European Space Agency by the Advanced Concepts Team that implements the differential algebra of Taylor truncated polynomials as well as Taylor models (which combine Taylor polynomials with a bounded interval), and a number of miscellaneous algorithms useful for its applications (differential intelligence, automatic differentiation, verified integration etc.).

Statement of need

pyaudi enables researchers to efficiently perform computations with high-order Taylor polynomials and Taylor models, which is relevant for a wide array of applications including differentiable intelligence, automatic differentiation, and verified integration applications.

Novel aspects

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The key novel aspects of this package are:

- Accelerated polynomial arithmetic to arbitrary dimension using Obake. Whereas other
 automatic differentiation packages typically are more suitable for quick simple systems,
 using Obake allows pyaudi to efficiently handle high-order derivatives and more complex
 systems.
- Effective Taylor models for verified integration of ODEs implemented using Bernstein polynomials for bounding the range of multivariate polynomials. The only other open-source package, called TaylorModels.jl, calculates bounds using Horner's scheme combined with interval arithmetic. A quick test in the next section shows significantl speedup for even a relatively simple trivariate polynomial.

28 Comparison with TaylorModels.il

- 29 To quickly test the performance of the Taylor model implementation in pyaudi against
- TaylorModels.jl, we take three functions; one univariate, one bivariate and one trivariate shown
- 31 below. We then construct Taylor models of all the variables separately and time the evaluation
- of the function value as a Taylor model.



$$\begin{split} f(x,y,z) &= \frac{4\tan(3y)}{3x + x\sqrt{\frac{6x}{-7(x-8)}}} - 120 - 2x - 7z(1+2y) \\ &- \sinh\left(0.5 + \frac{6y}{8y+7}\right) + \frac{(3y+13)^2}{3z} - 20z(2z-5) \\ &+ \frac{5x\tanh(0.9z)}{\sqrt{5y}} - 20y\sin(3z) \end{split}$$

$$g(x,y) = \sin(1.7x + 0.5)(y+2)\sin(1.5y)$$

$$h(x) = x(x-1.1)(x+2)(x+2.2)(x+2.5)(x+3)\sin(1.7x+0.5)$$

Dimen-	Package	Remainder Bound (Order 1)	Remainder Bound (Order 15)	Speed Comparison
h(x)	TaylorMod- els.jl	1e-15	1e-15	\sim 1.5–2 \times faster than pyaudi
	pyaudi	1e+2	1e-5	\sim 1.5–2 $ imes$ slower than TaylorModels.jl
g(x, y)	TaylorMod- els.jl	1e+1	1e-6	Slower: pyaudi is $5 \times$ faster (order 3), $15 \times$ faster (order 15), $7800 \times$ faster (order 1, edge case)
	pyaudi	1e+1	1e-6	Faster (see above)
f(x, y, z)	TaylorMod- els.jl	1e+0	1e-11	Slower: pyaudi is $8 \times$ faster (order 3), $155 \times$ faster (order 15), $13000 \times$ faster (order 1, edge case)
	pyaudi	1e-1	1e-17	Faster (see above)

In the table above, a clear trend can be seen both in terms of speed and accuracy. For univariate Taylor models, TaylorModels.jl is marginally faster and numerically precise. At two dimensions, the remainder bounds are of equal size, but pyaudi is significantly faster, with the speedup increasing with the order of the polynomial. At three dimensions, pyaudi produces significantly tighter bounds and is again significantly faster, with the speedup increasing with the order of the polynomial.

References

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- A number of references to relevant work and algorithms implemented in pyaudi are:
- (Biscani, 2020)
 - (Makino, 1998)
- (Titi & Garloff, 2019)
- Other software packages that do similar things are:
 - JAX (Bradbury et al., 2018)
 - TensorFlow (Abadi et al., 2015)
- PyTorch (Paszke et al., 2019)
- COSY INFINITY (Makino & Berz, 2006)



- DACE (Massari et al., 2018)
- TaylorSeries.jl/TaylorModels.jl (Benet et al., 2019)
 - CORA (Althoff, 2015)

52 Ongoing research

- EclipseNET (Acciarini, Biscani, et al., 2024) (Acciarini, Izzo, et al., 2025)
- CR3BP stochastic continuation (Acciarini, Baresi, et al., 2024)
- Long-term propagation (Caleb & Lizy-Destrez, 2020)
 - dSGP4 (Acciarini, Baydin, et al., 2025)

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