GRÖBNER BASIS ALGORITHMS FOR GRASSMAN ALGEBRAS IN A MAPLE PACKAGE

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Abstract

A brief discussion of an approach to calculating Gröbner bases in the Grassmann (exterior) algebra with emphasis on the ideal membership problem is presented. A new package written for computing Gröbner bases in Grassmann algebras for Maple 11 is introduced.

1 Introduction

This paper will give details of the algorithms used by the author for the Maple package TNB. The TNB package is capable of calculating Gröbner bases in the Grassmann algebra. Very little space is devoted to background information and more to the workings of the algorithms themselves.

Two separate types of Gröbner bases will be developed. However, only one will be capable of determining left ideal membership. The steps taken to arrive at each type of Gröbner basis will be similar to the steps taken to compute a

(*ii*) $_{i \ j} = _{j \ i}, 1 \quad i, j \quad m$ (*iii*) $e_{i \ j} = _{j}e_{i}, 1 \quad i \quad n, 1 \quad j \quad m$

Note that as a consequence of (i) above $e_i^2 = 0$. Thus there exist nonzero zero divisors. Also observe that $\bigwedge_{n,0}$ is a copy of the Grassmann algebra of order *n* over the field *k*. Note also that $\bigwedge_{0,m} = k[x_1, x_2, \ldots, x_m]$, is the ring of commutative polynomials in *m* variables over the field *k*. Thus it is not a big leap to see that the commutative techniques should produce similar results on the super Grassmann algebra as they do on $k[x_1, \ldots, x_n]$ since our commutative ring is embedded in the algebra. For the remainder of this paper, let m = 0, and so we will be working within the Grassmann algebra.

The quality of being Noetherian is a very important concept in the computation of commutative Gröbner bases as it guarantees termination of the division

- Deg[InvLex] order gives [*e*₁₂₃, *e*₂₃, *e*₁₃, *e*₁₂, *e*₃, *e*₂, *e*₁, 1]
- InvDeg[Lex] order gives [1, e₁, e₂, e₃, e₁₂, e₁₃, e₂₃, e₁₂₃]

Note that RevLex, InvRevLex, and InvDeg[Lex] orders are not admissible. The admissibility of the monomial order is important when redhTd[(e)]TJd; e

Algorithm 1 [4] Left Normal Form LeftNF(f/G)

1: h = f2: while h = 0 and $G_h = \{g \ G/LM(g)/LM(h)\} =$ do 3: choose $g \ G_h$ 4: h = Spoly(h, g)5: end while 6: return h

Theorem 1 (Characterization Theorem for GLBs). [7] The following conditions are equivalent.

- (i) F is a GLB.
- (ii) If $f_1, f_2 = F$, and $t = T_n$ satisfies $t \cdot \text{lcm}(\text{LMon}(f_1), \text{LMon}(f_2)) = 0$, then LeftNF $(t \cdot \text{LeftSpoly}(f_1, f_2)), F) = 0$.

Based on the characterizations in Theorem 1, Algorithm 2 will compute a GLB for a given list of Grassmann polynomials. Note that lines 6 and 7 of Algorithm 2 ensure that the "if" condition in (ii) of Theorem 1 is met while lines 8 through 17 perform the "then" statement of the Theorem.

Algorithm 2 [7] Gröbner left basis in Λ_n

1: G := F, $B = \{\{f_1, f_2\} | f_1, f_2 \quad G; f_1 = f_2\}$ 2: while B = do $\{f_1, f_2\}$:= an element of B 3: $B := B - \{f_1, f_2\}$ 4: $g := \text{LeftSpoly}(f_1, f_2)$ 5: $V := A - V(\operatorname{Icm}(\operatorname{LMon}(f_1), \operatorname{LMon}(f_2)))$ 6: $T := T(V) \{1\}$ 7: while T = do8: t := an element of T 9: $T := T - \{t\}$ 10: $h := t \cdot q$ 11: h := N(G, h)12: if h = 0 then 13: $B := B \{ \{g, h\} | g \in G \}$ 14: 15: $G := G \{h\}$ end if 16: end while 17. 18: end while 19: return G

De1926Tf21.536t1770/149922Td[13)-1(t343.711il)-1(e) T17Td[19143(oa[]-222(f)] Id1492Td[13)-1(3

Theorem 2 (Characterization Theorem for GLIBs). [7] The following statements are equivalent.

- (i) F is a GLIB.
- (ii) If $f_1, f_2 = F$, then for any t, LeftNF $(t \cdot f_1, F) = 0$ and LeftNF $(t \cdot LeftSpoly(f_1, f_2)$

3 The TNB Package

The TNB package is a package of procedures written for Maple 11 that can compute Gröbner left bases and Gröbner left ideal bases in a Grassmann algebra. When the package is loaded, 20 functions are exported.

> with(TNB);

TNB - package for computing Groebner bases in Grassmann algebras TNB version delta 0.3 (March 12, 2008) 20 functions exported

[Deg, InvDeg, InvLex, InvRevLex, LCoe, LMon, LTerm, Lex, NF, RevLex, TLSpolyG, TglbG, TglibG, TlcmG, TmingbG, idealpoly, isadmissible, makelist, pairs, testing]

Several procedures are from the package GFG [2]. These procedures are the monomial orderings and procedures that use the monomial orderings to find the leading term, monomial, and coe cients. The procedures are *Deg*, *InvDeg*, *InvLex*, *InvRevLex*, *LCoe*, *LMon*

a space of dimension *n*. The procedure does not return a zero polynomial. If one of its randomly generated polynomials happens to be zero, it recursively calls itself until it returns exactly *N* nonzero such polynomials. This procedure is useful for testing purposes.

- 9. Procedure *idealpoly* creates a random polynomial (Cli ord or Grassmann, depending on the second argument) in an one-sided ideal *J* left or right, depending on the third argument- and generated by polynomials supplied as the first list *F*. The second argument is expected to be wedge for the wedge product in a Grassmann algebra, or cmul for the Cli ord product in a Cli ord algebra. In either case, the algebra is considered over a space of dimension equal to the largest index found in the list *F*. An output of this procedure contains three items, in the following order:
 - (a) A list of random coe cients.
 - (b) A list of generators (it is a permuted list *F* due to removal of duplicates in the first line of the procedure when Maple may and usually does permute elements of *F*).
 - (c) Random polynomial. Assigning a name to this procedure will result in the random polynomial being assigned to that name.

This procedure is useful for testing purposes.

10. Procedure *testing* takes a list of polynomials F and for each t = T(V) and computes the product $t = f_i$ for i = 1, ..., n where n is the number of polynomials in the given list F then finds the LeftNF($t = f_i/F$). This is based on Theorem 2. It is useful in determining whether a given Gröbner basis is a GLIB.

Package TNB also relies upon the CLI FFORD package [1]. The CLI FFORD package provides the basis monomials and exterior, or wedge, product which is used for multiplication of Grassmann monomials. Type checking of arguments passed to TNB procedures is also provided by the CLI FFORD package. When loading, the TNB package automatically loads the CLI FFORD package and the BI GEBRA package which is part of the CLI FFORD library.

4 Applications and Examples

This section will show examples of the TNB package in use. In many examples the results are compared to those from

Example 2. [7] Given $f_1 = e_{56} - e_{13}$ and $f_2 = e_{45} - e_{23}$ procedures TglibG and

p4 := *e123*

P := [Id + 2e1 + 3e3 + e13, e1 + 3e23, e2 + e1, e123]First a GLIB will be computed by TNB. This Gröbner basis is capable of deter-

mining ideal membership.

> Tgb: =Tgl i bG(P, Deg[Lex]);

Computed 21 S-polynomials among 7 polynomials in Groebner basis and needed to compute 21 $\,$

This procedure took 3.495 seconds to run.

 $Tgb := [e123, \ Id + 2 \ e1 + 3 \ e3 + e13, \ \frac{1}{3}e1 + e23, \ e2 + e1, \ e2, \ e3, \ Id]$

P := [e1 + e3 + e13, e1 + e23, e2 + e1, e123]

> GBt: =Tgl i bG(P, Deg[Lex]);

Computed 15 S-polynomials among 6 polynomials in Groebner basis and needed to compute 15

This procedure took 3.494 seconds to run.

GBt := [e123, e1 + e3 + e13, e1 + e23, e2 + e1, -e3 + e2, e3]> GBtm: =Tmi ngbG(GBt, Deg[Lex]);

$$GBtm := [e2 + e1, -e3 + e2, e3]$$

The minimal Gröbner basis *GBtm* is capable of determining ideal membership. Now Plural will compute a reduced Gröbner basis.

> Pgb: =PLURALforGl i nk(P, 0, dp, [e1, e2, e3], i nput_for_Si ngul ar, i nput_for_Mapl e, ' i nfty', ' d');

$$Pgb := [e3, e2, e1]$$

> GBtm<>Pgb;

[e2 + e1, -e3 + e2, e3] = [e3, e2, e1]

Note that the lists GBtm and Pgb are not identical. The di erence is due to the fact that Gröbner basis returned by TNB is a minimal Gröbner basis and the one returned by Plural is a reduced Gröbner basis, which is a concept that is also applicable in the non-commutative case [5]. To show that they are in fact equal, it is su cient to show that each of the generators of one ideal is contained in the other. This can be accomplished by using the NF procedure to reduce each generator with respect to the generators of the other ideal. Since Pgb is a reduced Gröbner basis it can determine ideal membership.

> for i from 1 to 3 do NF(GBtm[i], Pgb, Deg[Lex]) end do;

0, 0, 0

This shows that every generator in the list GBtm is in the ideal generated by the polynomials (in this case monomials) of Pgb.

> for i from 1 to 3 do NF(Pgb[i], GBtm, Deg[Lex]) end do;

0,0,0

It is important to recall that a minimal Gröbner basis given by the procedure Tmi ngbG

> N: =4: numofpol s: =4: i:='i': P:=[]: for i from 1 to numofpols do p | | i : =0: end do: for i from 1 to numofpols do while p | |i=0 or p | |i=Id do p||i:=rd_clipolynom(N): end do: P: = [op(P), p||i];end do; vars: =[e||(seq(i,i=1..N))]; $P := [7 \ Id - 8 \ e14]$ $P := [7 \ Id - 8 \ e14, \ 6 \ Id + 6 \ e13]$ $P := [7 \ Id - 8 \ e^{14}, 6 \ Id + 6 \ e^{13}, -4 \ e^{1} + 5 \ e^{3}]$ P := [7 Id - 8e14, 6 Id + 6e13, -4e1 + 5e3, Id - 2e34 - 2e23 + e14 + e13 + e3]vars := [e1, e2, e3, e4]

Computed 15 S-polynomials among 6 polynomials in Groebner basis and needed to compute 15 $\,$

This procedure took 17.784 seconds to run.

$$GB1 := [Id - 2e34 - 2e23 + e14 + e13 + e3,$$
$$Id + e13, -\frac{7}{8}Id + e14, e1 - \frac{5}{4}e3, e3, Id]$$

> GB1m: =Tmi ngbG(TNBgl i b, Deg[Lex]);

$$Pgb := [Id]$$

> evalb(GB1m=Pgb);

true

The last result shows that the two lists are equivalent. Thus the ideal generated by the four polynomials is the whole algebra. Even though the polynomials were restricted to degree at most 4, any degree polynomial will be in the left ideal generated by them. This is from the fact that the reduced Gröbner basis is only the *Id* element and so any polynomial in any degree algebra will be a multiple of the generator.

Example 6. This example will use randomly generated polynomials of a higher degree. Instead of using four polynomials of degree up to 4, there will be two polynomials of unto degree 6 generated. That is the polynomials will be from the Grassmann algebra Λ_6 . The same procedure will be used as before.

> N: =6: numofpol s: =2: i:='i': P:=[]: for i from 1 to numofpols do p | | i : =0: end do: for i from 1 to numofpols do while p||i=0 or p||i=1d do p||i:=rd_clipolynom(N): end do: P: = [op(P), p | |i];end do; vars: =[e||(seq(i, i=1..N))]; P := [-e13 + 3 e46 - e6 - e2356]P := [-e13 + 3 e46 - e6 - e2356, e2]vars := [e1, e2, e3, e4, e5, e6]> GB1: =Tgl i bG(P, Deg[Lex]);

Computed 45 S-polynomials among 10 polynomials in Groebner basis and needed to compute 45 $\,$

This procedure took 16.769 seconds to run.

$$GB1 := [e1456 + \frac{1}{3}e156, e13 - 3e46 + e6 + e2356, e135 + 3e456 + e56, e146 - \frac{1}{3}e16, e156, e346 - \frac{1}{3}e36, e13 - 3e46 + e6, e16, e36, e2]$$

> GB1m: =Tmi ngbG(GB1, Deg[Lex]);

GB1m := [e13 - 3 e46 + e6, e16, e36, e2]
> Pgb: =PLURALforGl i nk(P, 0, dp, vars, i nput_for_Si ngul ar,
i nput_for_Mapl e, ' i nfty', ' d');

Pgb := [e2, e36, e16, e13 - 3 e46 + e6]
> eval b(convert(GB1m, set)=convert(Pgb, set));

true

In the last result the two lists GB1m and Pgb were converted to sets and compared since Maple usually puts lists in random order making it di cult to compare longer lists. The result is *true* and so the two sets are equal and thus the lists are equal. This means that the ideal generated by the polynomials that were randomly generated can be generated by the four polynomials given in GB1m and Pgb. Thus TNB gave the same result as PI ural did.

Example 7. This example will demonstrate that the GLB does not solve the membership problem but the GLIB will. A few extra lines of code also allow one to follow the computation of the GLB by listing out the S-polynomials as they are computed and the new polynomials as they are added to the GLB *G*.

> f1: =e2456-e3; f2: =e14-e36; f3: =e1256-e13; F: =[f1, f2, f3];