

HAPprime

**More homological algebra with prime
power groups**

Version 0.6

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Chapter 1

Introduction

1.1 Introduction to the HAPprime package

HAPprime is a package for the GAP computer algebra system (<http://www.gap-system.org/>), and which extends the HAP ‘Homological Algebra Programming’ package written by Graham Ellis (<http://hamilton.nuigalway.ie/Hap/www/>). It provides algorithms and data structures for calculating resolutions of small prime-power groups. The HAPprime functions use significantly less memory than the equivalent function in HAP, allowing resolutions (and cohomology ring presentations) of larger groups to be calculated (see Section 2.2).

Earlier versions of HAPprime also included functions to compute polynomial ring presentations of cohomology rings, and to ensure that these rings are complete and correct. These functions have now been moved into the HAP package and are documented as part of that package (see for example `Mod2CohomologyRingPresentation` (**HAP: Mod2CohomologyRingPresentation**) and `PoincareSeriesLHS` (**HAP: PoincareSeriesLHS (HAPprime)**)).

1.2 Required software

The HAPprime package requires GAP version 4.4 or greater and HAP version 1.9.3 or greater.

1.3 Installing HAPprime

To install the HAPprime Package, unpack the archive file into your GAP packages directory (either usually the `pkg` directory of your GAP 4 installation if you have access to it, or some local `pkg` directory that GAP can find). The HAPprime files will all be installed in a subdirectory called `happrime`.

1.4 Loading and testing HAPprime

The HAPprime package is not loaded by default when GAP is started. To load the package, type the following at the GAP prompt:

Example

```
gap> LoadPackage( "HAPprime");
```

If HAPprime isn't already in memory, it is loaded and the author information is displayed. If you are a frequent user of HAPprime, you might consider putting this line in your `.gaprc` file, or using the `PackagesToLoad` option in your `gap.ini` file.

The correct installation of HAPprime can be tested by using the test routine `tst/testall.g`:

Example

```
gap> ReadPackage("HAPprime", "tst/testall.g");
+ HAPprime version 0.6
  general tests
+ GAP4stones: 371057
+ HAPprime version 0.6
  userguide examples
+ GAP4stones: 387662
+ HAPprime version 0.6
  datatypes reference manual examples
+ GAP4stones: 382653
true
```

The number of GAP4stones will vary depending on your machine, but any additional lines of messages indicate problems with your installation.

The test routine calls a set of test files (**Reference: Test Files**) which can be found in the `tst` directory of the HAPprime installation. All of the routines listed in this user guide are tested, as are many of those in the datatype reference manual.

1.5 Documentation

The documentation for HAPprime is in two parts. This document is the user guide, which covers the main functions provided by HAPprime and examples of their use. There is also a more technical HAPprime datatypes reference manual which gives details of the new GAP datatypes defined and used internally by HAPprime, as well as outlining the algorithms used by the package.

1.5.1 MakeHAPprimeDoc

▷ `MakeHAPprimeDoc([manual-name])` (function)

Returns: nothing

The two manuals supplied with HAPprime - this user guide and the datatypes reference manual - are written using the `GAPDoc` package and are available in PDF, HTML and text format. It should not be necessary to rebuild these files, but should you wish to do so then this can be done using the function `MakeHAPprimeDoc`.

The optional argument *manual-name* is a string specifying which manuals to build. It may be one of the following

- "all" builds both manuals. This is the default
- "userguide" builds just the user guide
- "datatypes" builds just the datatypes reference manual
- "internal" builds both manuals, including the otherwise undocumented internal functions

- "testexamples" builds neither manual, but tests all of the examples using `TestManualExamples` (**GAPDoc: `TestManualExamples`**)

As well as building the manuals, this function at the same time builds **GAP** test files (**Reference: `Test Files`**) which means that all of the testable examples in the manuals are added to the HAPprime test routines described in Section 1.4.

1.6 Displaying progress and calculation information

By default, the functions in HAPprime display no output (except for returning the result). The `InfoHAPprime` info class can be used to enable the printing of progress and calculation information during processing. Since some computations with HAPprime can take several hours, setting this to a higher level can be particularly useful for monitoring the progress of computations.

1.6.1 InfoHAPprime

▷ `InfoHAPprime`

(info class)

The `InfoHAPprime` info class is used throughout the HAPprime package. Use `SetInfoLevel(InfoHAPprime, level)` to change the amount of information displayed about the progress of the computation (see `SetInfoLevel` (**Reference: `InfoLevel`**) in the **GAP** reference manual). The different distinct levels are:

- 0 print nothing (this is the default)
- 1 print some information, mainly progress information during computations that may take some time
- 2 print more detailed information, including details of internal calculations

Chapter 2

Examples

2.1 Computing the mod p group cohomology

Let G be a group and \mathbb{F} be a field, and let $\mathbb{F}G$ be the group ring of G over \mathbb{F} . A free $\mathbb{F}G$ -resolution of the ground ring is an exact sequence of module homomorphisms

$$\dots \rightarrow M_{n+1} \rightarrow M_n \rightarrow M_{n-1} \rightarrow \dots \rightarrow M_1 \rightarrow \mathbb{F}G \twoheadrightarrow \mathbb{F}$$

Where each M_n is a free $\mathbb{F}G$ -module and the image of $d_{n+1} : M_{n+1} \rightarrow M_n$ equals the kernel of $d_n : M_n \rightarrow M_{n-1}$ for all $n > 0$. The maps d_n are called boundary homomorphisms. In HAPprime we consider the case where G is a p -group and \mathbb{F} is the prime field $GF(p)$, and this is assumed from now on.

If we now define the Abelian group D_n to be $Hom(M_n, \mathbb{F})$, the set of all homomorphisms $M_n \rightarrow \mathbb{F}$, we can obtain the dual of this sequence, which will be a cochain complex of Abelian group homomorphisms

$$\dots \leftarrow D_{n+1} \leftarrow D_n \leftarrow D_{n-1} \leftarrow \dots \leftarrow D_1 \leftarrow \mathbb{F} \leftarrow \mathbb{F}$$

Each group D_n will be isomorphic to $\mathbb{F}^{|M_n|}$ where $|M_n|$ is the rank of the module M_n . Unlike the resolution, this sequence will generally not be exact, but one can define the mod- p cohomology group of G at degree n to be

$$H^n(G, \mathbb{F}) = \frac{\ker(D_n \rightarrow D_{n+1})}{\text{im}(D_{n-1} \rightarrow D_n)}$$

for all $n > 0$. As with the D_n , the mod- p cohomology groups will also be isomorphic to vector spaces over \mathbb{F} . In the case where the resolution R is minimal (where each module M_n has the minimal number of generators), the dimensions of the (co)homology groups $H^n(G, \mathbb{F})$ are identical to the dimensions of the resolution modules M_n . The group cohomology (and the similar group homology) is an invariant of G , and does not depend on a particular free $\mathbb{F}G$ -resolution.

In the general case, there are thus two stages to computing the group cohomology of G up to the n th cohomology group:

1. Compute R , a free $\mathbb{F}G$ -resolution for $\mathbb{F}G$, with at least $n + 1$ terms.
2. Construct the cochain complex C from R and compute the n homology groups of C

For example, to calculate the 9th mod- p cohomology group of the 134th order 64 in the GAP small groups library (which is the Sylow 2-subgroup of the Mathieu group M_{12}), we can use the HAPprime function `ResolutionPrimePowerGroupRadical` (3.1.1) to compute 10 terms of a free $\mathbb{F}G$ -resolution for G and then use HAP functions to find the rank b_9 of the cohomology group, which will be isomorphic to \mathbb{F}^{b_9} . Alternatively, since `ResolutionPrimePowerGroupRadical` (3.1.1) always returns a minimal resolution, the cohomology group dimensions can be read directly from the resolution.

— Example —

```
gap> G := SmallGroup(64, 134);;
gap> # First construct a FG-resolution for the group G
gap> R := ResolutionPrimePowerGroupRadical(G, 10);
Resolution of length 10 in characteristic 2 for <pc group of size 64 with
6 generators> .
No contracting homotopy available.
A partial contracting homotopy is available.

gap> # Convert this into a cochain complex (over the prime field with p=2)
gap> C := HomToIntegersModP(R, 2);
Cochain complex of length 10 in characteristic 2 .

gap> # And get the rank of the 9th cohomology group
gap> b9 := Cohomology(C, 9);
55
gap>
gap> # Since R is a free resolution, the ranks of the cohomology groups
gap> # are the same as the module ranks in R
gap> ResolutionModuleRanks(R);
[ 3, 6, 10, 15, 21, 28, 36, 45, 55, 66 ]
```

2.2 Comparing the memory usage and speed of HAPprime and HAP's ResolutionPrimePowerGroup functions

For small p -groups, the group ring $\mathbb{F}G$ can be considered as a vector space of rank $|G|$ with the elements of G as its basis elements. Each module M_n in a $\mathbb{F}G$ -resolution is also a vector space (of dimension $|M_n||G|$) and the boundary maps d_n can be represented as vector space homomorphisms. As a result, standard linear algebra techniques can be used to compute a minimal resolution by constructing a sequence of module homomorphisms where the kernel of one map is the image of the next, and where the modules have minimal generating sets. See Chapter (**HAPprime Datatypes: Resolution construction functions**) in the datatypes manual for further details.

As the groups get larger, this approach becomes less feasible due to the amount of time and memory needed to store and compute the null space of large matrices. The HAP function `ResolutionPrimePowerGroup` (**HAP: ResolutionPrimePowerGroup**) and the HAPprime functions `ResolutionPrimePowerGroupRadical` (3.1.1) and `ResolutionPrimePowerGroupGF` (3.1.1) all use this linear algebra approach, but the HAPprime functions are optimised to save memory, allowing the computation of resolutions which are longer, or are of larger groups, than are possible using HAP alone.

2.2.1 HAPprime takes less memory to store resolutions

Consider computing a resolution of a group of an arbitrary group of order 128, $G = \text{SmallGroup}(128, 844)$ using HAP. Computation is performed on a dual-core Intel Core2Duo running at 2.66MHz, and the memory available to GAP is the standard initial allocation of 256Mb.

Example

```
gap> G := SmallGroup(128, 844);;
gap> R := ResolutionPrimePowerGroup(G, 9);
Resolution of length 9 in characteristic 2 for <pc group of size 128 with
7 generators> .

gap> time;
27685
gap> # Can we construct a resolution of length ten?
gap> R := ResolutionPrimePowerGroup(G, 10);
exceeded the permitted memory ('-o' command line option) at
res := SemiEchelonMatDestructive( List( mat, ShallowCopy ) );
called from
SemiEchelonMat( NullspaceMat( BndMat ) ) called from
ZGbasisOfKernel( i - 1 ) called from
<function>( <arguments> ) called from read-eval-loop
Entering break read-eval-print loop ...
you can 'quit;' to quit to outer loop, or
you can 'return;' to continue
```

The HAPprime function `ResolutionPrimePowerGroupRadical` (3.1.1) uses an almost identical algorithm, but stores its boundary maps more efficiently. As a result, with the same memory allowance:

Example

```
gap> G := SmallGroup(128, 844);;
gap> R := ResolutionPrimePowerGroupRadical(G, 9);
Resolution of length 9 in characteristic 2 for <pc group of size 128 with
7 generators> .
No contracting homotopy available.
A partial contracting homotopy is available.

gap> time;
25321
gap> # Can we construct a resolution of length ten?
gap> R := ExtendResolutionPrimePowerGroupRadical(R);;
gap> # Yes! How about eleven?
gap> R := ExtendResolutionPrimePowerGroupRadical(R);
Resolution of length 11 in characteristic 2 for <pc group of size 128 with
7 generators> .
No contracting homotopy available.
A partial contracting homotopy is available.

gap> ResolutionModuleRanks(R);
[ 3, 6, 11, 19, 30, 44, 62, 85, 113, 146, 185 ]
gap>
gap> # But it will run out of memory if we try to go to twelve terms
gap> R := ExtendResolutionPrimePowerGroupRadical(R);
exceeded the permitted memory ('-o' command line option) at
...
```

The HAPprime version can compute two further terms of the resolution, which given the sizes of the additional modules represents a considerable improvement. Just representing the homomorphism $d_{10} : (\mathbb{F}G)^{146} \rightarrow (\mathbb{F}G)^{113}$ as vectors requires nearly as much memory again as representing the first nine homomorphisms. To compute and store the same resolution of length 11 using `ResolutionPrimePowerGroup` (**HAP: ResolutionPrimePowerGroup**) would need a little over three times the memory used here by HAPprime. The time taken by both versions is very similar.

In the example above, note also the use of the HAPprime function `ExtendResolutionPrimePowerGroupRadical` (3.1.2), which makes it much easier to add terms to an existing resolution. In standard HAP, if one decides that a resolution is too short and that more terms are required, then the entire resolution must be computed again from scratch.

2.2.2 HAPprime takes less memory to compute resolutions

The function `ResolutionPrimePowerGroupGF` (3.1.1) uses a new algorithm to compute the kernel of $\mathbb{F}G$ -module homomorphisms when $\mathbb{F}G$ -modules are represented using a set of G -generating vectors (see (**HAPprime Datatypes: FG-module homomorphisms**) in the datatypes reference manual). This provides a further memory saving over `ResolutionPrimePowerGroupRadical` (3.1.1), although at the cost of a much slower computation time:

Example

```
gap> G := SmallGroup(128, 844);;
gap> R := ResolutionPrimePowerGroupGF(G, 9);
Resolution of length 9 in characteristic 2 for <pc group of size 128 with
7 generators> .
No contracting homotopy available.
A partial contracting homotopy is available.

gap> time;
422742
gap> R := ExtendResolutionPrimePowerGroupGF(R);;
gap> R := ExtendResolutionPrimePowerGroupGF(R);;
gap> R := ExtendResolutionPrimePowerGroupGF(R);;
gap> R := ExtendResolutionPrimePowerGroupGF(R);;
gap> R := ExtendResolutionPrimePowerGroupGF(R);;
gap> R := ExtendResolutionPrimePowerGroupGF(R);;
Resolution of length 15 in characteristic 2 for <pc group of size 128 with
7 generators> .
No contracting homotopy available.
A partial contracting homotopy is available.

gap> ResolutionModuleRanks(R);
[ 3, 6, 11, 19, 30, 44, 62, 85, 113, 146, 185, 231, 284, 344, 412 ]
gap> # But it will run out of (the initial 256Mb) of memory at sixteen terms
```

Using `ResolutionPrimePowerGroupGF` (3.1.1) we can get a further four terms of the resolution. For this resolution, this represents a memory saving of a factor of five over `ResolutionPrimePowerGroupRadical` (3.1.1) and fifteen over `ResolutionPrimePowerGroup` (**HAP: ResolutionPrimePowerGroup**), although it does take fifteen times as long as either of those just to compute the first nine terms, and scales less well with size.

2.2.3 Automatic selection of the best method

The two functions `ResolutionPrimePowerGroupRadical` (3.1.1) and `ResolutionPrimePowerGroupGF` (3.1.1) offer a trade-off between time and memory. The function `ResolutionPrimePowerGroupAutoMem` (3.1.1) automates the decision of which version to use, switching from the Radical to the GF version when it estimates that it is about to run out of available memory for the faster version. In this example, we have also increase the `InfoHAPprime` (1.6.1) info level to display progress information. At level two, the rank of each module in the resolution is displayed as it is calculated, giving an indication of progress. With this setting, the user is also notified when the `AutoMem` function switches, and the GF function displays a rolling estimate of its completion time (which is not shown since that output is overwritten when completed)

Example

```
gap> G := SmallGroup(128, 844);;
gap> SetInfoLevel(InfoHAPprime, 2);
gap> R := ResolutionPrimePowerGroupAutoMem(G, 15);
#I Dimension 2: rank 6
#I Dimension 3: rank 11
#I Dimension 4: rank 19
#I Dimension 5: rank 30
#I Dimension 6: rank 44
#I Dimension 7: rank 62
#I Dimension 8: rank 85
#I Dimension 9: rank 113
#I Finding kernel of homomorphism by splitting:
#I   - Finding kernel of U
#I   - Finding kernel of V
#I   - Finding intersection of U and V
#I   - Finding intersection preimages
#I Dimension 10: rank 146
#I Finding kernel of homomorphism by splitting:
#I   - Finding kernel of U
#I   - Finding kernel of V
#I   - Finding intersection of U and V
#I   - Finding intersection preimages
#I Dimension 11: rank 185
#I Finding kernel of homomorphism by splitting:
#I   - Finding kernel of U
#I   - Finding kernel of V
#I   - Finding intersection of U and V
#I   - Finding intersection preimages
#I Dimension 12: rank 231
#I Finding kernel of homomorphism by splitting:
#I   - Finding kernel of U
#I   - Finding kernel of V
#I   - Finding intersection of U and V
#I   - Finding intersection preimages
#I Dimension 13: rank 284
#I Finding kernel of homomorphism by splitting:
#I   - Finding kernel of U
#I   - Finding kernel of V
#I   - Finding intersection of U and V
#I   - Finding intersection preimages
```

```
#I Dimension 14: rank 344
#I Finding kernel of homomorphism by splitting:
#I   - Finding kernel of U
#I   - Finding kernel of V
#I   - Finding intersection of U and V
#I   - Finding intersection preimages
#I Dimension 15: rank 412
Resolution of length 15 in characteristic 2 for <pc group of size 128 with
7 generators> .
No contracting homotopy available.
A partial contracting homotopy is available.

gap> StringTime(time);
" 5:45:53.613"
```

Chapter 3

Functions for Homological Algebra

3.1 Resolutions

3.1.1 ResolutionPrimePowerGroup

- ▷ `ResolutionPrimePowerGroupRadical(G , n)` (operation)
- ▷ `ResolutionPrimePowerGroupGF(G , n)` (operation)
- ▷ `ResolutionPrimePowerGroupAutoMem(G , n)` (operation)
- ▷ `ResolutionPrimePowerGroupGF2(G , n)` (operation)
- ▷ `ResolutionPrimePowerGroupRadical(M , n)` (operation)
- ▷ `ResolutionPrimePowerGroupGF(M , n)` (operation)
- ▷ `ResolutionPrimePowerGroupAutoMem(M , n)` (operation)
- ▷ `ResolutionPrimePowerGroupGF2(M , n)` (operation)

Returns: `HAPResolution`

Returns n terms of a minimal free $\mathbb{F}G$ -resolution for either the ground ring of a prime power group G or of a module M . For the module version, M must be passed as an `FpGModuleGF` object - see (**HAPprime Datatypes: FG-modules**) in the HAPprime datatypes reference manual.

Three versions of this function are provided:

ResolutionPrimePowerGroupRadical

uses the same resolution-building method as the HAP function `ResolutionPrimePowerGroup` (**HAP: ResolutionPrimePowerGroup**), but stores the resolution in a different format that takes only about half the memory of the HAP version.

ResolutionPrimePowerGroupGF

calculates the resolution using HAPprime's G -generator form of modules, which reduces memory use by around a factor of two over `ResolutionPrimePowerGroupRadical`, but is slower by an order of magnitude.

ResolutionPrimePowerGroupAutoMem

automatically switches between the two previous versions based on the available memory. It uses the Radical version until it gets close to the limit of the available memory, and then switches to the GF version.

ResolutionPrimePowerGroupGF2

calculates the resolution by $\mathbb{F}G$ -matrix partitioning. The amount of partitioning is

governed by the (**Reference: Options Stack**) option `MaxFGExpansionSize`. The default value means that until the boundary map takes about 128Mb, the method is equivalent to `ResolutionPrimePowerGroupRadical`, and then it tends towards `ResolutionPrimePowerGroupGF` in terms of time, but saves less memory.

See the HAPprime datatypes reference manual for details of the different algorithms, in particular the chapters on the G -generator form of $\mathbb{F}G$ -modules (**HAPprime Datatypes: FG-modules**) and $\mathbb{F}G$ -module homomorphisms (**HAPprime Datatypes: FG-module homomorphisms**) and on resolutions (**HAPprime Datatypes: Resolutions**).

3.1.2 ExtendResolutionPrimePowerGroup

- ▷ `ExtendResolutionPrimePowerGroupRadical(R)` (operation)
- ▷ `ExtendResolutionPrimePowerGroupGF(R)` (operation)
- ▷ `ExtendResolutionPrimePowerGroupAutoMem(R)` (operation)
- ▷ `ExtendResolutionPrimePowerGroupGF2(R)` (operation)

Returns: `HAPResolution`

Returns the resolution R extended by one term. The three variants offer a choice between memory and speed, and correspond to the different versions of `ResolutionPrimePowerGroup` in HAPprime. See the documentation (3.1.1) for those functions for a description of the different variants.

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